SEQUENCE LISTING

<110> Kleanthous, Harold

Al-Garawi, Amal Miller, Charles Tomb, Jean Francois Oomen, Raymond P. <120> Identification of Polynucleotides Encoding Novel Helicobacter Polypeptides in the Helicobacter Genome <130> 06132/043002 <140> US 09/895,913 <141> 2001-06-29 <150> TS 08/881,227 <151> 1997-06-24 <160> 368 <170> FastSEQ for Windows Version 4.0 <210> 1 <211> 1004 <212> DNA <213> Helicobacter pylori <220> <221> CDS <222> (56)...(931) cattaaacgc atgatttttg ctacaataat aggattttaa ttatataaag gacaa atg ggc atg cca aat agg ggc gtt gtt tta tta gac ggg caa gcg cta gct Gly Met Pro Asn Arg Gly Val Val Leu Leu Asp Gly Gln Ala Leu Ala gat aat ata gaa aaa gat ttg aaa cat aaa atc caa ata ata acc gca Asp Asn Ile Glu Lys Asp Leu Lys His Lys Ile Gln Ile Ile Thr Ala 154 25 caa acg cat aaa cgc ccc aaa cta gcc gtg att tta gtg ggg aaa gat Gln Thr His Lys Arg Pro Lys Leu Ala Val Ile Leu Val Gly Lys Asp 202 45 ccc gct agt atc act tat gtc aat atg aag atc aaa gca tgc gaa agg Pro Ala Ser Ile Thr Tyr Val Asn Met Lys Ile Lys Ala Cys Glu Arg 250 55

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	aa ttg ys Lei													346
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	cc cct hr Pro	-												538
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gtc to Val Se 2														922

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Val Cys His Ile Leu Thr Lys Asp Ile Ser Phe Tyr Thr Gln Asn Ala
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					gga Gly											1406
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Phe Ile Leu Tyr Thr			Iis Ala Leu		Val
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Tyr Phe Ser Ile Ala	Ile Ile I				Val
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Leu His Glu Asp Thr	Leu Thr I	120 Ile Leu L		125 Gly Glu Tyr	Pro
130 Ile Phe Ser Ser Phe		Phe Leu I		Val Leu Thr	
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2 2 · · · · · · · · · · · · · · · · · ·		-	_ = Don		-1~

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			Asp	645					650					655	
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                         25
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														aat Asn		1496
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Phe Asn Glu Asp Lys Ala Ile Leu Thr Thr Glu Asn Gln Gly Leu Tyr
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Leu Glu Gly Val Asn Ser Phe Asp Glu Val Glu Arg Ser Arg Leu Lys
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Thr Ser Arg Ala Lys Asn Tyr Tyr Val Ser Val Ala Arg Arg Gly Ala
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	cgc Arg 100														392
	ccc Pro														440
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				ttg Leu										1928
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Ile Leu Ile Asn Ala Asn Ser Tyr Ser Val Ala Asp Val Glu Lys Ile

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			Glu 260					265					270		
		275	Glu				280					285			
	290		Val			295					300				
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Val	Leu	Val	Val 340	Phe	Ala	Phe	Leu	Arg 345	Asn	Gly	Thr	Ile	Thr 350	Leu	Val
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_	-	-		_	_	act Thr			_			_	2600
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Val Ser Ala Phe Lys Asn Ala Ile Met Glu Ile Leu Asn Ser Lys Thr
                                985
Val Gly Asp Val Phe Gly Glu Asn Gly Leu Leu Asn Ala Leu Asp Pro
                            1000
                                               1005
Thr Glu Arg Lys Lys Ile Asp Gln Met Leu Leu Glu Gln Ile Gln Ala
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                                            1020
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			1460					1469					1470		
Lys	Arg	Met 1475		Tyr	Gly	Tyr	Asp 1480		Gln	Ile	Ile	Arg 1489		Gly	Ser
Leu	Ser 1490	_	Tyr	Leu	Lys	Leu 149		Thr	Leu	Ile	Asp 150	Phe 0	Asn	Gly	Lys
Arg 150		Gln	Leu	Asn	Gly 1510	_	Ser	Leu	Ser	Tyr 151		Asn	Gln		Val 1520
Asn	Ile	Lys	Asp	Gly 152		Leu	Val	Val	Ser 153		Lys	Asp	Asn	Gln 1535	
Gl'n	Met	Val	Tyr 1540		Ser	Ile	Leu	Tyr 154		Lys	Val	Gln	Val 1550		Val
Ser	Asp	Lys 1555		Met	Asp	Ile	His 1560		Pro	Ser	Leu	Glu 1565	_	Tyr	Ile
Lys	Tyr 1570		Gln	Gly	Ser	Ala 157	_	Leu	Asp	Ala	Ile 158	·Lys	Ser	Ala	Gly
Asn 1585		Ser	Ile	Leu	Trp 1590		Asn	Glu	Leu	Phe 1595		Ala	Lys	_	Gly L600
<u>Asn</u>	Pro	Fen	Phe	Ala 160		፲ንェ	Tyr	Leu	Gln 161	_	Asn	bro	Thr	Glu 1615	
Ile	Val	Thr	Leu 1620		Lys	Asp	Ile	Thr 1625		Ala	Leu	Gly	Met 1630		Ser
Lys	Pro	Asn 1635		Lys	Asn	Asn	Ser 1640		Asp	Ala	Leu	Gln 1645		Asn	Thr
Tyr	Thr 1650		Gln	Met	Ser	Arg 1655		Ala	Lys	Leu	Ser 1660	Asn)	Phe	Ala	Ser
Phe 1665		Ser	Thr	Asp	Phe 1670		Glu	Arg	Leu	Ser 1675		Leu	Lys		G1n L680
Arg	Phe	Ala	Asp	Ala 1689		Pro	Asn	Ala	Met 1690		Val	Ile	Leu	Lys 1699	_
Ser	Gln	Arg	Asp 1700		Leu	Lys	Asn	Asn 1705		Trp	Ala	Thr	Gly 1710		Gly
Gly	Val	Ser 1715		Val	Glu	Asn	Gly 1720		Gly	Thr	Leu	Tyr 1725		Val	Asn
Val	Gly 1730		Asp	Arg	Phe	Ile 1735		Gly	Val	Ile	Val 1740	Gly)	Gly	Tyr	Ala
Ala 1745		Gly	Tyr	Ser	Gly 1750		Tyr	Glu	Arg	Ile 1755		Asn	Ser	-	Ser 1760
				1765	5				1770)		Ile	_	1775	5
Glu	Leu	Thr	Phe 1780	Ser	Val	Asn	Glu	Thr 1785		Gly	Ala	Asn	Lys 1790		Gln
Ile	Ser	Ser 1795		Asp	Thr	Leu	Leu 1800		Met	Ile	Asn	Gln 1805		Tyr	Lys
Tyr	Ser 1810		Trp	Thr	Thr	Asn 1815		Lys	Val	Asn	Tyr 1820	Gly)	Tyr	Asp	Phe
Met 1825		Lys	Asn	Lys	Ser 1830		Ile	Leu	Lys	Pro 1835		Ile	Gly		Arg .840
Tyr	Tyr	Tyr	Ile	Gly 1845		Thr	Gly	Leu	Glu 1850		Val	Met	His	Asn 1855	
Leu	Tyr	Asn	Gln 1860		Lys	Ala	Asn	Ala 1865		Pro	Ser	Lys	Lys 1870		Val
Leu	Thr	Ile 1875		Leu	Ala	Leu	Glu 1880		Arg	His	Tyr	Phe 1885		Thr	Asn
Ser	Tyr 1890		Tyr	Ala	Ile	Gly 1895		Phe	Gly	Arg	Asp 1900	Leu)	Leu	Val	Asn

1905 1910 1915 1920 Tyr Arg Lys Gly Glu Leu Tyr Asn Thr Phe Ala Ser Ile Thr Thr Gly 1925 1930 Gly Glu Val Arg Leu Phe Lys Ser Phe Tyr Ala Asn Ala Gly Val Gly 1945 Ala Arg Phe Gly Leu Asp Tyr Lys Met Ile Asn Ile Thr Gly Asn Ile 1955 1960 1965 Gly Met Arg Leu Ala Phe 1970 <210> 13 <211> 755 <212> DNA <213> Helicobacter pylori <220> <221> CDS <222> (124)...(690) <400> 13 ttgcgcttat aaattcccgg ctttagaaaa acacaccaaa attgtaggag tcattaacca 60 agtggggggc assgggcgat cacaccggtc gctcttttag agcctgtgga aattgctgga 120 gct atg att aat aga gcg acc tta cac aat tat tct gaa att gaa aaa Met Ile Asn Arg Ala Thr Leu His Asn Tyr Ser Glu Ile Glu Lys aag aat atc atg ctc agt gat agg gtc gtt gtc att aga agc ggc gat 216 Lys Asn Ile Met Leu Ser Asp Arg Val Val Ile Arg Ser Gly Asp 25 30 gtg atc cct aaa atc atc aag cct tta gaa tct tat aga gac ggc tcg 264 Val Ile Pro Lys Ile Ile Lys Pro Leu Glu Ser Tyr Arg Asp Gly Ser 35 40 caa cat aaa att gaa cgc ccc aag gtt tgc cct ata tgt tcg cat gag 312 Gln His Lys Ile Glu Arg Pro Lys Val Cys Pro Ile Cys Ser His Glu 50 ctt ttg tgc gaa gag att ttt act tat tgt caa aac ctt aat tgc ccg 360 Leu Leu Cys Glu Glu Ile Phe Thr Tyr Cys Gln Asn Leu Asn Cys Pro 65 70 gca agg ttg aaa gaa agc ttg att cat ttc gct tct aaa gac gct tta 408 Ala Arg Leu Lys Glu Ser Leu Ile His Phe Ala Ser Lys Asp Ala Leu 80 85 90 aac att caa ggc ttg ggc gat aaa gtc ata gag caa ctt ttt gaa gaa 456 Asn Ile Gln Gly Leu Gly Asp Lys Val Ile Glu Gln Leu Phe Glu Glu 100 105 aag ctc att ttt aac gct ctg gat ttg tat gct tta aaa tta gaa gat 504 Lys Leu Ile Phe Asn Ala Leu Asp Leu Tyr Ala Leu Lys Leu Glu Asp 115 120

Ser Met Gly Asp Lys Leu Val Arg Phe Ile Gly Asn Asn Thr Leu Ser

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gac gct att tta aaa agc aaa aac cct ccc tta tgg cgt ttg att aac
                                                                   600
Asp Ala Ile Leu Lys Ser Lys Asn Pro Pro Leu Trp Arg Leu Ile Asn
    145
                        150
gct tta ggg att gag cat att ggt aag gga gcg agt aaa acg ctg gcc
                                                                   648
Ala Leu Gly Ile Glu His Ile Gly Lys Gly Ala Ser Lys Thr Leu Ala
                    165
                                        170
                                                                   690
aaa tac ggc tta aat gtg tta gaa aaa agc gaa scg agt ttt
Lys Tyr Gly Leu Asn Val Leu Glu Lys Ser Glu Xaa Ser Phe
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His Lys Ile Glu Arg Pro Lys Val Cys Pro Ile Cys Ser His Glu Leu
                        55
                                            60
Leu Cys Glu Glu Ile Phe Thr Tyr Cys Gln Asn Leu Asn Cys Pro Ala
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Arg Leu Lys Glu Ser Leu Ile His Phe Ala Ser Lys Asp Ala Leu Asn
                                    90
Ile Gln Gly Leu Gly Asp Lys Val Ile Glu Gln Leu Phe Glu Glu Lys
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Leu Ile Phe Asn Ala Leu Asp Leu Tyr Ala Leu Lys Leu Glu Asp Leu
        115
                            120
                                                125
Met Arg Leu Asp Lys Phe Lys Ile Lys Lys Ala Gln Asn Leu Leu Asp
                        135
                                            140
Ala Ile Leu Lys Ser Lys Asn Pro Pro Leu Trp Arg Leu Ile Asn Ala
                    150
                                        155
Leu Gly Ile Glu His Ile Gly Lys Gly Ala Ser Lys Thr Leu Ala Lys
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Tyr Gly Leu Asn Val Leu Glu Lys Ser Glu Xaa Ser Phe
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atg cgg ccg ggt gtt ttg caa gtg ggg gcg atg tat gct aat ggg gta
                                                                    104
Met Arg Pro Gly Val Leu Gln Val Gly Ala Met Tyr Ala Asn Gly Val
                             10
ggg atc caa acc aac aga tta aaa gcc gct cgc tat tat gaa tgg gtt
                                                                    152
Gly Ile Gln Thr Asn Arg Leu Lys Ala Ala Arg Tyr Tyr Glu Trp Val
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                         25
                                              30
                                                                    200
gca gcg gcg gcg atg cga ccg ctt gcg cga atc tgg ctc aga tgt atg
Ala Ala Gly Ala Met Arg Pro Leu Ala Arg Ile Trp Leu Arg Cys Met
 35
                     40
aaa aca aga aaa atg cgg att caa acg ata aag aaa acg ctt tgc aat
                                                                    248
Lys Thr Arg Lys Met Arg Ile Gln Thr Ile Lys Lys Thr Leu Cys Asn
                 55
                                                          65
                                                                   296
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Cys Met Arg Trp Leu Val Lys Gly Gly Ile Cys Ser His Ala Ile Ile
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                                  75
                                                                   344
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Trp Gly Gly Cys Leu Leu Thr Glu Val Gly Ser Gln Lys Ile Ile Thr
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Lys Arg
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Gly Val Gly Ile Gln Thr Asn Arg Leu Lys Ala Ala Arg Tyr Tyr Glu
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Trp Val Ala Ala Gly Ala Met Arg Pro Leu Ala Arg Ile Trp Leu Arg Cys Met Lys Thr Arg Lys Met Arg Ile Gln Thr Ile Lys Lys Thr Leu 55 Cys Asn Cys Met Arg Trp Leu Val Lys Gly Gly Ile Cys Ser His Ala Ile Ile Trp Gly Gly Cys Leu Leu Thr Glu Val Gly Ser Gln Lys Ile Ile Thr Lys Arg 100 <210> 17 <211> 1837 <212> DNA <213> Helicobacter pylori <220> <221> CDS <222> (51)...(1784) <400> 17 cgcatgcgct cctttctaaa gcgatcaaaa acaaagagta agggattaac atg tca 56 Met Ser aaa aaa atc gta gtc gat cct atc act agg att gag ggg cat tta agg 104 Lys Lys Ile Val Val Asp Pro Ile Thr Arg Ile Glu Gly His Leu Arg 10 att gaa gtg atc gta gat gat gat aac gtg atc act gat gcg ttt tct 152 Ile Glu Val Ile Val Asp Asp Asp Asn Val Ile Thr Asp Ala Phe Ser 25 30 tct tct acg ctt ttt agg ggg cta gaa acc att att aaa ggc aga gat 200 Ser Ser Thr Leu Phe Arg Gly Leu Glu Thr Ile Ile Lys Gly Arg Asp 35 40 cca cga gat gca ggc ttc atc gct caa agg att tgc ggg gta tgc act 248 Pro Arg Asp Ala Gly Phe Ile Ala Gln Arg Ile Cys Gly Val Cys Thr 55 65 tat tcg cat tat aag gcc ggt atc acg gcg gta gaa aac gct cta ggc 296 Tyr Ser His Tyr Lys Ala Gly Ile Thr Ala Val Glu Asn Ala Leu Gly 70 atc act ccc cca tta aac gcg caa ttg gtg cga tct ttg atg aac atg 344 Ile Thr Pro Pro Leu Asn Ala Gln Leu Val Arg Ser Leu Met Asn Met 85 90 gcg ctg ctt ttt cat gac cat gtg gtg cat ttc tat act ttg cat ggg 392 Ala Leu Leu Phe His Asp His Val Val His Phe Tyr Thr Leu His Gly 105 ctt gat tgg tgc gat atc atg agc gct tta aaa gcc gat ccc att caa 440

Leu 115	Asp	Trp	Cys	Asp	Ile 120	Met	Ser	Ala	Leu	Lys 125	Ala	Asp	Pro	Ile	Gln 130	
	_				ttc Phe											488
_					gcg Ala											536
					cct Pro											584
					gag Glu											632
_			-		caa Gln 200		_				_		_			680
					cac His								-	-	Thr	728
					ttg Leu											776
-		_			gcc Ala						_				_	824
		_	_		gaa Glu	_		_		-			_			872
					aat Asn 280											920
					ttg Leu											968
					att Ile											1016
cat His	tct Ser	tgg Trp 325	tat Tyr	caa Gln	tac Tyr	gaa Glu	gac Asp 330	act Thr	aaa Lys	gaa Glu	gtg Val	caa Gln 335	ctc Leu	cac His	cct Pro	1064

								tta Leu 350					1112
_								aaa Lys					1160
						_	-	tac Tyr	_	_	-		1208
_	-	-			_	_	 _	ggt Gly					1256
								tta Leu					1304
			-				_	cga Arg 430					1352
								ctt Leu	_			_	1400
								act Thr					1448
								tac Tyr					1496
								aaa Lys					1544
								aat Asn 510					1592
								agc Ser	_				1640
								att Ile					1688
								gtg Val					1736

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ggg cag tct tta aac gag ttt aaa gta gag cct aat ttc gct aaa ttc

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Gly Glu Ser Val Gly Ile Glu Asn Lys Ile Ile Pro Ala Lys Val Leu
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                                                 365
Asp Thr Lys Asn Lys Tyr Ser Trp Ile Lys Ser Pro Arg Tyr Asp Ser
                        375
Lys Pro Met Glu Val Gly Pro Leu Ser Ser Val Val Val Gly Leu Ala
                    390
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Ala Lys Asn Pro Tyr Val Thr Glu Val Ala Thr Lys Phe Leu Lys Asp
                405
                                    410
Thr Lys Leu Pro Leu Glu Ala Leu Phe Ser Thr Leu Gly Arg Thr Ala
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            420
                                                     430
Ala Arg Cys Ile Glu Ala Lys Thr Ile Ala Asp Asn Gly Leu Leu Ala
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                            440
                                                 445
Phe Asp Ala Leu Val Glu Asn Leu Lys Ser Asp Gln Ser Thr Cys Ala
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                                             460
Pro Tyr His Ile Asp Lys Asn Gln Glu Tyr Lys Gly Arg Tyr Ile Gly
                    470
                                        475
Gln Val Pro Arg Gly Met Leu Ser His Trp Val Arg Ile Lys Asn Gly
                485
                                    490
Val Val Glu Asn Tyr Gln Ala Val Val Pro Ser Thr Trp Asn Ala Gly
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                                                     510
Pro Arg Asp Ser Gln Asn Gln Arg Gly Ala Tyr Glu Met Ser Leu Ile
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                                                 525
Gly Thr Lys Ile Ala Asp Leu Thr Gln Pro Leu Glu Ile Ile Arg Thr
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                                            540
Ile His Ser Phe Asp Pro Cys Ile Ala Cys Ser Val His Val Met Asp
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                     Met Ser Gln Lys Ile Leu Ile Leu Gly Ile Gly
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aat atc ctt ttt ggc gat gaa ggg att ggg gtg cat tta gcc cac tac
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Asn Ile Leu Phe Gly Asp Glu Gly Ile Gly Val His Leu Ala His Tyr
             15
ctc aaa aaa aat ttt tct ttt ttc cct agc gtg gat att ata gat ggg
                                                                   208
Leu Lys Lys Asn Phe Ser Phe Phe Pro Ser Val Asp Ile Ile Asp Gly
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		a ggc gtt gag ata gga u Gly Val Glu Ile Gly 70	
		t aaa gaa atc aca tgg o Lys Glu Ile Thr Try 5 90	Ala
		c act tta agg ctc acg s Thr Leu Arg Leu Thi 105	
		c gtg ggg ctt gtg cct e Val Gly Leu Val Pro 120	
		t tca agc aaa att tta u Ser Ser Lys Ile Leu 135	
Ala Leu Glu Thr Ala		a acc caa ctc aac gca u Thr Gln Leu Asn Ala 150	
		c gct tta gaa tgt ato e Ala Leu Glu Cys Ile 5 170	e Ala
gaa ctt tct tat aag Glu Leu Ser Tyr Lys 175		tttgtttttc tttttaaat	.g 643
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20	25	s Tyr Leu Lys Lys Asr 30	
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Gln Leu Ile Pro Leu : 50	Ile Thr Ser Tyr Glu 55	ı Lys Val Leu Ile Leu 60	ı Asp

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Cys Val Ser Ala Glu Gly Val Glu Ile Gly Ser Val Tyr Ala Phe Asp

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392
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                        105
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Asn Tyr Gln Gln Lys Ile Ala Asn Ala Thr Gly Leu Ala Leu Lys Asn
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Pro Pro Leu Lys Gln Asn Ile Pro Asn Lys Ser Leu Ser Leu Glu Asn
                            40
Leu Asn Glu Gln Ile Lys Asn Leu Lys Ser Val Pro Ala Leu Glu Lys
                        55
                                             60
Leu Lys Asn Asp Phe Asn Glu Arg Phe Lys Ile Cys Pro Lys Asp Lys
Glu Thr Leu Trp Leu Glu Thr Ile Gln Lys Leu Val Ala Ser Glu Phe
                                    90
Phe Glu Leu Glu Asp Ala Ile Asn Phe Gly Gln Glu Leu Glu Asn Ala
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											gcg Ala		152
											gtc Val		200
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											agc Ser		296
_	_		_	_	_					_	 att Ile		344
_	_			_	_						gaa Glu		392
		-		_		-		_	_		ttg Leu		440
											aaa Lys 145		488
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											atc Ile		584
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											cta Leu 225		728

		aaa Lys														776
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_		gtg Val	_													872
		tct Ser														920
		act Thr														968
		gat Asp														1016
		gca Ala 325					-		-		_	_	_			1064
		aag Lys														1112
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Lys Phe Asn Lys Pro Phe Glu Leu Val Leu Glu Asp Met Leu Lys Ala
                    150
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Gly Val Asp Ser Met Pro Gly Gly Gly Ala Glu Ile Phe Asp Glu Glu
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Ile Arg Arg Lys Ile Cys Asn Gly Lys Val Gly Ser Ser Arg Trp Leu
            180
                                185
Glu Ile His Ala Tyr Trp His Lys Leu Gly Lys Met Ser Asn Ala Thr
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                                                 205
Met Leu Phe Gly His Ile Glu Asn Lys Ile His Arg Ile Asp His Met
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                                            220
Leu Arg Ile Lys Lys Ile Gln Ser Pro Lys Asn Gln Val Glu Asn Lys
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Glu Gly Gly Phe Asn Ala Phe Ile Pro Leu Leu Tyr Gln Lys Glu Asn
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Asn Tyr Leu Asn Val Glu Lys Ser Pro Ser Ala Ile Glu Ile Leu Lys
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Thr Ile Ala Ile Ser Arg Ile Leu Leu Asn Asn Ile Pro His Ile Lys
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Ala Tyr Trp Ala Thr Leu Gly Leu Asn Leu Ala Leu Val Ala Gln Glu
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Phe Gly Ala Asn Asp Leu Asp Gly Thr Ile Glu Ile Glu Ser Ile Gln
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Ser Ala Ala Gly Ala Lys Ser Arg His Gly Leu Glu Lys Glu Asp Leu
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Lys Leu Pro Lys Lys Arg Val Ser Lys Thr Lys Ser Gln Lys Leu Ile
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cca gat aa Pro Asp As			Leu A	-	_	_	_	_		296
gaa gcc ag Glu Ala Ar 8										344
ccg att ga Pro Ile As 100										392
aat tta cg Asn Leu Ar 115				_						440
ttt gtc aa Phe Val Ly										488
gac aac at Asp Asn Me	_		Pro Ph					_		536
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Leu Ile Hi	20		25	5	·		30			
Ile Ser Al 35		Met Leu	Leu Gl	ıu Leu	Glu Lys	Gly 45	Ala	Phe	Lys	

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Lys Asn Glu Ala Tyr Phe Ile Ser Asp Glu Glu Asp Lys Asn Tyr Val

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cgc aa Arg Ly: 115														440
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aaa gg Lys Gl	_			_				_		_				536
caa aa Gln Ly		Phe						_	_	_			_	584
ccc at Pro Ile 18	e Gln													632
tac cag Tyr Gli 195	-			_	_	_	_				_			680
att caa Ile Gl														728
ttt acc	_					-				-				776
gaa tgg Glu Trp														824
gaa gcc Glu Ala 260	a Thr													872
tgg gcg Trp Ala 275								tgaa	agcgo	ett g	gaaa	aaat	c	919
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His Leu Pro Ile Asp Thr Leu Ile Glu Lys Ala Arg Glu Lys Ile Ala
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                                    90
Gln Lys His Gly Ile Ala Trp Tyr Lys Lys Ser Tyr Phe Ser Leu Leu
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Lys Asn Arg Lys Lys Met Asp Arg Val Phe Ser Thr Ala His Phe Leu
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Val Phe Lys Gly Leu Phe Lys Arg Phe Lys Lys Ser Ala Leu Pro Pro
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Leu Asn Gln Lys Ser Phe Leu Gln Lys His Ala Glu Met Lys Leu Leu
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Glu Asn Pro Ile Gln Lys Val Ala Ile Phe Ile Gly Cys Leu Ser Asn
                                185
Tyr His Tyr Gln Gln Val Gly Glu Ser Leu Leu Tyr Ile Leu Glu Lys
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                                            220
Ala Tyr Phe Thr Gly Asp Lys Asp Thr Thr Leu Phe Leu Val Lys Lys
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Asn Ile Glu Trp Phe Glu Ser Tyr Leu Asp Lys Val Asp Ala Ile Ile
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-52-

205

729

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Arg Ile As		Arg A	Asp I		Gly 235	Lys	Asp	Lys	Asn	Thr 240	Thr	Arg	Gln	
gaa tac ga Glu Tyr As 245			Asn S											825
att tca aa Ile Ser Ly 260		Lys I												873
aat tct ct Asn Ser Le														921
caa ata ca Gln Ile Hi							_							969
tat ata ga Tyr Ile Gl 31	lu Asn													1017
gcc cac co Ala His Pr 325			3ln 7		_	_		_		_				1065
atc tgc tt Ile Cys Le 340		Ala I												1113
gta tat ga Val Tyr As						-		_		_				1161
aca act at Thr Thr Il									-				-	1209
ata gat go Ile Asp Al	la Phe			Asp										1257
tca tac tt Ser Tyr Le 405			ys A											1305
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Thr Asn Ile Glu Asp Phe Glu Lys His Leu Glu Gln Ser Lys Asp Met
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Gly Lys Glu Ile Asn Tyr Leu Ser Val Cys Thr Pro Thr His Thr His
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                                 75
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                                                                   346
Phe Asp His Ile Arg Phe Gly Leu Arg Asn Gly Met His Val Ile Cys
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                             90
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Glu Lys Pro Leu Val Leu Asp Pro Gly Glu Ile Gln Glu Leu Lys Asp
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                        105
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115
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                                        125
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Leu His Cys Asp Thr Leu Ala Leu Lys Glu Lys Ile Lys Ser Glu Leu
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gac aaa aac cct agc aag gtg ttt gac atc acg ctc act tat atc agc
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Asp Lys Asn Pro Ser Lys Val Phe Asp Ile Thr Leu Thr Tyr Ile Ser
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                                155
gtt caa ggg aaa tgg tat ttt tct tca tgg cga gcg gat gtg aat agg
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Tyr Leu Phe Gly Ser Val Lys Asp Lys Val Ile Asn Lys Glu Glu 195 200 205 210 205 210 210 205 210 210 205 210 210 205 210 210 215 210 215 215 215 215 215 215 219 2212 PRT 2213 Helicobacter pylori 2400 > 32 Met Leu Phe Ala Met Ile Gly Ser Gly Gly Phe Ile Ala Pro Lys His 1	729
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Leu Gln Ala Ile Arg Asp Thr Gly His Phe Leu Asp Cys Ser Phe Asp 20	
Solution Solution	
50	
70 75 80 Thr His Phe Asp His Ile Arg Phe Gly Leu Arg Asn Gly Met His Val 85 90 95 Ile Cys Glu Lys Pro Leu Val Leu Asp Pro Gly Glu Ile Gln Glu Leu	
85 90 95 Ile Cys Glu Lys Pro Leu Val Leu Asp Pro Gly Glu Ile Gln Glu Leu	
Lys Asp Leu Glu Val Lys His Gln Lys Arg Val Phe Ser Leu Leu Pro 115 120 125	
Leu Arg Leu His Cys Asp Thr Leu Ala Leu Lys Glu Lys Ile Lys Ser 130 135 140	
Glu Leu Asp Lys Asn Pro Ser Lys Val Phe Asp Ile Thr Leu Thr Tyr 145 150 155 160	
Ile Ser Val Gln Gly Lys Trp Tyr Phe Ser Ser Trp Arg Ala Asp Val 165 170 175	
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					cct Pro											632
					atc Ile 200			_						_	-	680
					aaa Lys											728
					aaa Lys	_		-							_	776
	_				ttt Phe	_	-		_	-	_					824
					ttt Phe					_						· 872
			_		aac Asn 280		-		_							920
_		-			acc Thr			_						~	-	968
	_	_	_		gct Ala				_						~	1016
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Phe 355	Val	Leu	Met	Thr	ttt Phe 360	Leu	Ser	Ala	Gly	Ile 365	Val	Asn	Ile	Phe	Ile 370	1160
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					gcg Ala											1352
					att Ile 440											1400
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                                            220
Lys Leu Leu Lys Asp Glu Tyr Lys Glu Ile Glu Leu Ile Ser His Gln
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Gln Asp Lys Thr Ile Ala His Phe Leu Glu Asn Ser Ala Leu Leu Ser
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Tyr Ala Gly Ile Met Gly Met Met Ala Ser His Ser Val Gly Gly His
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Lys Thr Phe Val Leu Met Thr Phe Leu Ser Ala Gly Ile Val Asn Ile
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Leu Pro Ala Gly Gln Ser Leu Gly Val Asp Pro Gly Val Val Ser Met
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                                        395
Ala Ile Ala Trp Gly Asp Ala Trp Thr Asn Met Ile Gln Pro Phe Trp
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165

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	_		_	_		_				_		_		att Ile		200
				_	_	_						_	_	gtt Val 65		248
			_			_								aat Asn		296
		_	-		_		_			_		-		ctt Leu		344
		_										_		gat Asp	_	392
_				-						-	-		_	tta Leu		440
_		_			_	-	_		_					aac Asn 145	_	488
	_	_					_				_	_	_	act Thr		536
														atg Met		584
_		-											_	agt Ser		632
												_	-	ttt Phe		680
														aac Asn 225		728

											ttg Leu					776
											gtc Val					824
											ttg Leu 270					872
											aaa Lys					920
-		_		_	_		_		-		caa Gln		_			968
											aac Asn					1016
											ata Ile					1061
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Thr Asn Ala Glu Asn Ile Leu Ala Tyr Thr Phe Phe Gln Glu Gly Leu
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Met Glu Ala Gln Glu Asn Leu Ser Leu Lys Asn Ile Phe Lys Thr Leu
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                                                     190
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Ser Ser Lys Asp Ala Lys Ala Ile Glu Glu Leu Gln Asp Lys Leu Arg
                            200
Phe Ser Ala Pro Lys Leu Gly Val Ser Ile Gln Ala His His Leu Lys
                        215
                                             220
Asn Leu Leu Glu Ala Phe Tyr His Gln Asn Lys Glu Ser Leu Gly Phe
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                                         235
Phe Ser Pro Tyr Phe Ser Leu Arg Ser Gln Thr Pro Ser Val Ser Tyr
                245
                                    250
Glu Ser Ala Leu Ala Ser Leu Glu Asn Tyr Phe Met Ala Leu Phe Gln
                                265
            260
Ser His Phe Lys Asp Asp Thr Ala Leu Gln Gln Asn Phe Lys Gly Leu
                            280
Leu Gln Ala Phe Val Ser Met Ala Lys Asp Lys Arg Ser Gln Ile Ala
                        295
                                             300
Leu Asn Ala Gln Ala Lys Asp Asn Ala Lys Leu Thr Phe Asn Ala Leu
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Leu Glu Ser Leu Ser Val Asn Phe Phe Gln Ser Tyr Lys Ile Ser His
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                                                                   104
Ile Lys Thr His Ser Ser Asn Glu Lys Glu Arg Phe Val Arg Ile Glu
gag gac gaa aag aaa gga tta ttt gct gga act gca aat gaa aat tcg
                                                                   152
Glu Asp Glu Lys Lys Gly Leu Phe Ala Gly Thr Ala Asn Glu Asn Ser
     20
                         25
                                              30
cac ggc ctt tct tta atg gct tta ata ggg gta ttg gtt ttt ggg ggc
                                                                   200
His Gly Leu Ser Leu Met Ala Leu Ile Gly Val Leu Val Phe Gly Gly
 35
                     40
gcg ttt tta gct ctg tta gcg cct aaa atc tat tta agc aat aat atc
                                                                   248
Ala Phe Leu Ala Leu Leu Ala Pro Lys Ile Tyr Leu Ser Asn Asn Ile
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145

296

tat tat att agc cgt aaa atc aac acc cta gaa gat caa aaa cgc ctg Tyr Tyr Ile Ser Arg Lys Ile Asn Thr Leu Glu Asp Gln Lys Arg Leu 70 75 ctt tta gaa gag caa caa atc cta aaa aac gaa tta gaa aaa gag cgt 344 Leu Leu Glu Glu Gln Gln Ile Leu Lys Asn Glu Leu Glu Lys Glu Arg 85 90 ttt aaa tac tac ata gaa aat agt gaa aat att ggc gat att gcg ttt 392 Phe Lys Tyr Tyr Ile Glu Asn Ser Glu Asn Ile Gly Asp Ile Ala Phe 100 105 110 taagtgaaaa accccctatc cccttaagag agcttaatta atggtcaatc att 445 <210> 38 <211> 114 <212> PRT <213> Helicobacter pylori <400> 38 Met Asn Ile Lys Thr His Ser Ser Asn Glu Lys Glu Arg Phe Val Arg 1.0 Ile Glu Glu Asp Glu Lys Lys Gly Leu Phe Ala Gly Thr Ala Asn Glu 25 Asn Ser His Gly Leu Ser Leu Met Ala Leu Ile Gly Val Leu Val Phe Gly Gly Ala Phe Leu Ala Leu Leu Ala Pro Lys Ile Tyr Leu Ser Asn Asn Ile Tyr Tyr Ile Ser Arg Lys Ile Asn Thr Leu Glu Asp Gln Lys 75 Arg Leu Leu Glu Glu Gln Gln Ile Leu Lys Asn Glu Leu Glu Lys 90 Glu Arg Phe Lys Tyr Tyr Ile Glu Asn Ser Glu Asn Ile Gly Asp Ile 100 105 Ala Phe <210> 39 <211> 218 <212> DNA <213> Helicobacter pylori <220> <221> CDS <222> (52)...(165) <400> 39 aaggttttta gcgaaataag ccccacgcc aaacacgccc atgctcatta g cac aat 57 His Asn

atc gca cat aaa gca caa agc gca aat caa aaa cac ata att cct agc

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Ile Ala His Lys Ala Gln Ser Ala Asn Gln Lys His Ile Ile Pro Ser
         5
                              10
                                                  15
cat ccc cct ttc cac aat aaa caa gga ttg cgc ccc cac cgc cgc aca
                                                                    153
His Pro Pro Phe His Asn Lys Gln Gly Leu Arg Pro His Arg Arg Thr
     20
                         25
caa aga aat cgc taaaccaaaa ccttctataa aaaccacaaa catcttgctt
                                                                    205
Gln Arg Asn Arg
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aatcctttca ctc
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<213> Helicobacter pylori
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Pro Ser His Pro Pro Phe His Asn Lys Gln Gly Leu Arg Pro His Arg
Arg Thr Gln Arg Asn Arg
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                                                         Met Ser
ggc att tta gag cct tta gga gca aat tac ttg acc ctt tta atg gtt
                                                                    104
Gly Ile Leu Glu Pro Leu Gly Ala Asn Tyr Leu Thr Leu Leu Met Val
         5
                             10
                                                  15
tgg atc cct acg act ttt tta gca tgc atg ctc acg gca ttt att atg
                                                                    152
Trp Ile Pro Thr Thr Phe Leu Ala Cys Met Leu Thr Ala Phe Ile Met
     20
                         25
                                              30
ggt ttt act gat ttg aaa tta gac agc gat ccg cat tat tta gag cgc
                                                                   200
Gly Phe Thr Asp Leu Lys Leu Asp Ser Asp Pro His Tyr Leu Glu Arg
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35					40					45					50	
														aaa Lys 65		248
				_										ggg Gly		296
														gct Ala		344
														ttc Phe		392
	_									-				gct Ala		440
_					-						-			tgc Cys 145	-	488
	-						_		_				_	aat Asn		536
					_		_			_		_	_	tat Tyr	_	584
														tat Tyr		632
														gcc Ala		680
		_	_			_			_					gct Ala 225	_	728
			-	_		_								act Thr		776
														ggc Gly		824
tat	gtg	ttt	gat	cat	gcg	ttt	ttg	atc	cct	ggg	gtt	tta	gtc	gtg	tct	872

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Tyr Val Phe Asp His Ala Phe Leu Ile Pro Gly Val Leu Val Val Ser
    260
                        265
                                            270
                                                                   914
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Leu Ser Val Ala Leu Gly Phe Val Val Ala Pro Leu Val Leu
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                           40
Glu Arg Leu Lys Ala Gly Lys Ile Ser Pro Pro Lys Ile Lys Glu Glu
                        55
                                            60
Lys Glu Thr Ser Lys Asn Ala Lys Leu Ser Leu Trp Ile Phe Ile Gly
                    70
                                        75
Gly Val Val Ala Ile Val Phe Tyr Ala Ser Ala Ile Ser Lys Asn Ile
                                    90
Ala Phe Val Ser Pro Val Val Leu Gly Arg Asp His Ala Ile Val Ser
                                105
Phe Met Leu Ser Val Ala Thr Leu Ile Val Leu Phe Cys Lys Ile Asn
                            120
Ala Asn Glu Ile Ala His Ser Ser Val Phe Lys Ser Gly Met Gln Ala
                        135
                                            140
Cys Val Cys Val Leu Gly Val Ala Trp Leu Gly Asp Thr Phe Val Ser
                    150
                                        155
Asn His Ile Asp Glu Ile Lys Arg Tyr Ala Ser Phe Leu Ile Ala Asp
                165
                                    170
Tyr Pro Phe Leu Leu Ala Val Ala Leu Phe Leu Ala Ser Met Leu Leu
                                185
Tyr Ser Gln Ala Ala Thr Ser Lys Ala Leu Ile Pro Ser Val Ile Thr
                            200
                                                205
Ala Leu Gly Ile Ser Ala Asn His Thr Glu His Leu Tyr Ile Ile Val
                        215
                                            220
Ala Ser Phe Ala Ser Val Ser Ala Leu Phe Val Leu Pro Thr Tyr Pro
                    230
                                        235
Thr Leu Leu Gly Ala Ile Ala Met Asp Asn Thr Gly Thr Thr Lys Met
                245
                                    250
Gly Arg Tyr Val Phe Asp His Ala Phe Leu Ile Pro Gly Val Leu Val
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Val Ser Leu Ser Val Ala Leu Gly Phe Val Val Ala Pro Leu Val Leu
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175

170

gct aat ggc g Ala Asn Gly Va 180		: Ile Ala Gln			632							
ata gcg ggg t Ile Ala Gly Le 195					680							
cta tac aaa aa Leu Tyr Lys Ly					728							
aat tac gcc to Asn Tyr Ala So 23	~		-		776							
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tta gcc gcc t Leu Ala Ala Pl 260		ı Ala Ile Ile			872							
aag ttt aaa ga Lys Phe Lys G 275					920							
gcg ttt atc go Ala Phe Ile A					968							
ttg caa aaa gt Leu Gln Lys Va 33	al His Ala Ile		Val Asn Ala		1016							
gta gtc caa gg Val Val Gln Gl 325					1064							
ctt ttt atc ac Leu Phe Ile Th 340		Gly Thr Ser			1112							
atc gct gtg tt Ile Ala Val Pb 355					1160							
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Met Ile Ser Ala Thr Leu Ile Ala Gly Leu Met Gly Gly Leu Gly Ile
                            40
Thr Glu Ser Phe Asn Ala Met Ile Asp Gly Met Lys Gly Asn Leu Asn
                        55
Ile Ala Leu Ser Tyr Ile Leu Leu Gly Ala Leu Ala Val Ala Ile Ala
                    70
Lys Ser Asn Leu Ile Lys Val Ala Leu Ser Lys Leu Ile Gly Leu Met
                                    90
                85
Asp Tyr Lys Arg Ser Thr Phe Cys Phe Leu Ile Ala Phe Ile Ala Cys
                                105
Phe Ser Gln Asn Leu Val Pro Val His Ile Ala Phe Ile Pro Ile Leu
                            120
                                                125
Ile Pro Pro Leu Leu His Leu Met Asn Arg Leu Glu Leu Asp Arg Arg
                        135
                                            140
Ala Val Ala Cys Ala Leu Thr Phe Gly Leu Gln Ala Pro Tyr Leu Val
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                                        155
Leu Pro Val Gly Phe Gly Leu Ile Phe Gln Thr Thr Ile Leu Glu Gln
                165
                                    170
Leu Lys Ala Asn Gly Val Ser Thr Thr Ile Ala Gln Ile Thr Gly Val
                                185
Met Trp Ile Ala Gly Leu Ala Met Val Val Gly Leu Leu Val Ala Val
        195
                            200
Leu Thr Leu Tyr Lys Lys Pro Arg His Tyr Lys Glu Lys Ser Phe Asn
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Ile Glu Asn Tyr Ala Ser Leu Gln Leu Asn Tyr His Asp Tyr Leu Thr
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                                        235
Phe Ile Gly Ile Val Val Ala Phe Val Ile Gln Leu Ala Thr Asp Ser
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                                    250
Met Pro Leu Ala Ala Phe Leu Ala Leu Ala Ile Ile Leu Leu Gly Arg
           260
                                265
Gly Ile Lys Phe Lys Glu Thr Asp Ser Leu Met Asp Asp Ser Val Lys
                            280
Met Met Ala Phe Ile Ala Phe Val Met Leu Val Ala Ser Gly Phe Gly
                        295
Glu Val Leu Gln Lys Val His Ala Ile Glu Gly Leu Val Asn Ala Ile
                    310
                                        315
Thr Ser Val Val Gln Gly Lys Leu Leu Gly Ala Phe Leu Met Leu Val
                325
                                    330
Val Gly Leu Phe Ile Thr Met Gly Ile Gly Thr Ser Phe Gly Thr Ile
                                345
                                                    350
Pro Ile Ile Ala Val Phe Tyr Val Pro Leu Cys Ala Lys Leu Gly Phe
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Ser Val Glu Ser Thr Ile Leu Leu Ile Ala
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Val Gly Leu Val Thr Thr Leu Ala Leu Gly Ile Ser Cys Ile Ile
tta ttc gtg gcg ttt ttt agc atc att tgg cgg att ttg gct ttg att
                                                                   156
Leu Phe Val Ala Phe Phe Ser Ile Ile Trp Arg Ile Leu Ala Leu Ile
             20
                                 25
tct ggg ttc cct tct tgc tgt ttg tat tcg tcc tta act tct tgt tta
                                                                   204
Ser Gly Phe Pro Ser Cys Cys Leu Tyr Ser Ser Leu Thr Ser Cys Leu
         35
                             40
gtc att ttt aaa gag ttg gtg tat tgg cgg cgt ttg atc gct aaa tct
                                                                   252
Val Ile Phe Lys Glu Leu Val Tyr Trp Arg Arg Leu Ile Ala Lys Ser
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                         55
ata aaa gcc aag aca aaa aat aaa aat aaa agc gaa gaa atg agc cat
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Ile Lys Ala Lys Thr Lys Asn Lys Asn Lys Ser Glu Glu Met Ser His
 65
                     70
aac gcc tta ttt tta aac cac aac tgg cct tgt aaa ttc aaa aga
                                                                   348
Asn Ala Leu Phe Leu Asn His Asn Asn Trp Pro Cys Lys Phe Lys Arg
gcc gca tgg ttt aat tcc cct aaa aac aaa gaa aag atg aaa aac ccc
                                                                   396
Ala Ala Trp Phe Asn Ser Pro Lys Asn Lys Glu Lys Met Lys Asn Pro
                                105
aga aaa aaa gct aaa aaa act ttt aag gtg atc aaa ctc cca tca agg
                                                                   444
Arg Lys Lys Ala Lys Lys Thr Phe Lys Val Ile Lys Leu Pro Ser Arg
                            120
age ttt ttt aaa gaa aaa agg ttt ttg acg cca ttg ata ggg ttg att
                                                                   492
Ser Phe Phe Lys Glu Lys Arg Phe Leu Thr Pro Leu Ile Gly Leu Ile
    130
                        135
                                            140
tta gaa aat tta ggc tca atg act tta ggg gca aag agc cag cca aat
                                                                   540
Leu Glu Asn Leu Gly Ser Met Thr Leu Gly Ala Lys Ser Gln Pro Asn
145
                  . 150
                                        155
                                                             160
tgc aag aca tta gat aaa aac gcc acc act aaa atg att aaa atc
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Cys Lys Thr Leu Asp Lys Asn Ala Thr Thr Thr Lys Met Ile Lys Ile
                                    170
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636
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Gly Lys Ser Asn Lys Lys Val Ser Leu Ala Ser Trp Leu Asn Ser Ser
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ctaaagccat ccaccacca tataaaaaaa acaaaaatac taattagccc ggccaataac 756
cccaaaaccc ccaccacttc catgetetta ggcacattgc cttettetet ggctttttgg 816
atttttttcg cgctagggag ttcggttttt tcttcttcag ccattggccc tcttttttaa 876
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Ser Gly Phe Pro Ser Cys Cys Leu Tyr Ser Ser Leu Thr Ser Cys Leu
                            40
Val Ile Phe Lys Glu Leu Val Tyr Trp Arg Arg Leu Ile Ala Lys Ser
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Ile Lys Ala Lys Thr Lys Asn Lys Asn Lys Ser Glu Glu Met Ser His
                    70
Asn Ala Leu Phe Leu Asn His Asn Asn Trp Pro Cys Lys Phe Lys Arg
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                                    90
Ala Ala Trp Phe Asn Ser Pro Lys Asn Lys Glu Lys Met Lys Asn Pro
                                105
                                                     110
Arg Lys Lys Ala Lys Lys Thr Phe Lys Val Ile Lys Leu Pro Ser Arg
                            120
                                                 125
Ser Phe Phe Lys Glu Lys Arg Phe Leu Thr Pro Leu Ile Gly Leu Ile
                                            140
                        135
Leu Glu Asn Leu Gly Ser Met Thr Leu Gly Ala Lys Ser Gln Pro Asn
                    150
                                        155
Cys Lys Thr Leu Asp Lys Asn Ala Thr Thr Thr Lys Met Ile Lys Ile
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                165
Gly Lys Ser Asn Lys Lys Val Ser Leu Ala Ser Trp Leu Asn Ser Ser
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											ctc Leu					297
_	-	_	_				~				tac Tyr 95	_				345
											ttc Phe					393
											agc Ser					441
											gtt Val		-			489
											tgc Cys					537
											gtt Val 175					585
											acg Thr					633
											atg Met					681
aaa	gaa	tac	tat	ggc	atg	cga	ttg	ggt	ttt	tat	ttt	tta	ggc	ctg	tta	729

Lys	Glu	Tyr	Tyr 215	Gly	Met	Arg	Leu	Gly 220	Phe	Tyr	Phe	Leu	Gly 225	Leu	Leu	
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	att Ile 245	taaa	agcto	caa a	atago	gaata	ag ct	caaag	gcctt	t ttg	gatto	gagt	gttt	tttt	tag	833
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1				5					10					15		
Leu	Phe	Phe	Leu 20	Leu	Phe	Leu	Phe	Asn 25	Val	Glu	Thr	Leu	Leu 30	Leu	Ser	
His	Phe	Ser 35	Asp	Asp	Phe	Ser	Gln 40	Leu	Phe	Phe	Leu	Phe 45	Glu	Asn	His	
Val	Tyr 50		Phe	Ile	Val	Lys 55		Asp	Tyr	Leu	Gly 60		Ile	Gly	Val	
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Trp	Asn	Phe	Pro 100	Val	Lys	Asp	Ser	Leu 105	Met	Val	Leu	Tyr	Leu 110	Phe	Tyr	
Phe	Ala	Leu 115	Leu	Ala	Thr	Leu	Leu 120	Trp	Arg	Phe	Leu	Gly 125	Ala	Ser	Met	
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Ala 145	Ser	Ser	Leu	Gln	Ser 150	Phe	Arg	Phe	Leu	Ser 155	Val	Ser	Asp	Cys	Val 160	
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Glu Ala Ala Leu Lys Thr Asn Leu Lys Arg Gln Asp Ile Leu Asn Tyr

170	1	75	180	
		ta ggc aaa aag gag eu Gly Lys Lys Glu 195		691
		ta ttc aaa cgc att eu Phe Lys Arg Ile 210		739
cgt ttt tagtgat Arg Phe	tcg ctgagaatgt	aggcttaaaa ttcaga	aagg gtgtttttaa	795 ·
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Glu Ser Phe Leu 35	Phe Arg Arg I	le Asp Ala Gly Phe 0	Ile Ser Ser Ile 45	
Ala Gly Tyr Pro 50	Phe Ala Leu H: 55	is Ser His Ser Leu 60	Gly Ile Val Ala	
Tyr Lys Glu Val 65	Leu Ser Val Le	eu Val Val Asp Thr 75	Lys Asn Ala Phe 80	
Asp Lys Glu Ser	Ala Ser Ser As 85	sn Ala Leu Ser Gln 90	Ala Leu Gly Leu 95	
Lys Gly Glu Val 100	Leu Ile Gly As	sn Lys Ala Leu Gln 105	Phe Tyr Tyr Ser 110	
Asn Pro Lys Lys 115		sp Leu Ala Ala Leu 20	Trp Tyr Glu Lys 125	
Lys Arg Leu Pro 130	Phe Val Phe Gl	ly Arg Leu Cys Tyr 140	Tyr Gln Asn Lys	
Asp Phe Tyr Lys 145	Arg Leu Ser Le	eu Ala Phe Lys His 155	Gln Lys Thr Lys 160	
Ile Pro Tyr Tyr	Ile Leu Lys Gl	lu Ala Ala Leu Lys 170	Thr Asn Leu Lys 175	
Arg Gln Asp Ile 180	Leu Asn Tyr Le	eu Gln Lys Ile Tyr 185		
Lys Lys Glu Gln 195		ys Ala Phe Tyr Arg 00		
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Leu Leu Phe Leu Glu Ala Glu Asp Pro Glu Lys Asp Ile Gly Leu Tyr
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Ile Asn Ser Pro Gly Gly Val Ile Thr Ser Gly Leu Ser Ile Tyr Asp
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                                        75
Thr Met Asn Phe Ile Arg Pro Asp Val Ser Thr Ile Cys Ile Gly Gln
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Ala Ala Ser Met Gly Ala Phe Leu Leu Ser Cys Gly Ala Lys Gly Lys
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Arg Phe Ser Leu Pro His Ser Arg Ile Met Ile His Gln Pro Leu Gly.
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                                                 125
Gly Ala Gln Gly Gln Ala Ser Asp Ile Glu Ile Ile Ser Asn Glu Ile
                        135
                                            140
Leu Arg Leu Lys Gly Leu Met Asn Ser Ile Leu Ala Gln Asn Ser Gly
                    150
                                        155
Gln Ser Leu Glu Gln Ile Ala Lys Asp Thr Asp Arg Asp Phe Tyr Met
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Ser Ala Lys Glu Ala Lys Glu Tyr Gly Leu Ile Asp Lys Val Leu Gln
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			-	aaa Lys	_				-						144
				aaa Lys											192
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				caa Gln											288
_				tac Tyr	_	_				-	_	_	_	_	336
				caa Gln											384
				ttg Leu	_		_								432
				gat Asp 150											480
				ttg Leu									_		528
				gct Ala	_					_	_			_	576
				aat Asn											624
				tgg Trp											672
				atg Met 230											720

		_		-	gcg Ala											768
		_			aaa Lys									_	_	816
			-		ctc Leu					-		tag	tttt	tta		862
aat	tttt	gaa a	aact	agaa	tt aa	aaac	cgcti	tt!	tataa	aact	gg					904
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Lys	Cys 50	Leu	Arg	Gly	Lys	Thr 55	His	Pro	Pro	Cys	Phe 60		Lys	Pro	Lys	
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Gly	Thr	Ser	Val	Val 85	Gln	Met	Ser	Trp	Leu 90	Gln	Ser	Arg	Glu	Lys 95	Phe	
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Ile	Tyr	Gly 115	Tyr	Lys	Gln	Phe	Phe 120	Pro	Lys	Lys	Glu	Arg 125	Tyr	Gly	Phe	
Arg	Phe 130	Tyr	Val	Ser	Leu	Asp 135	Tyr	Ala	Tyr	Gly	Phe 140	Phe	Leu	Lys	Asn	
Lys 145	Gly	Val	Leu	Gly	Asp 150		Leu	Arg	Glu	Ser 155		Gln	Ile	Pro	Lys 160	
	Tyr	Arg	Glu	Lys 165	Leu	Gln	Arg	Lys	Glu 170		Phe	Ile	Asn	Ala 175		
Phe	Tyr	Gly	Ala 180		Ala	Asp	Phe	Leu 185		Lys	Arg	Ala	Phe 190		Thr	
Leu	Ile	Leu 195		Met	Asn	Phe	Val 200		Glu	Thr	Trp	Phe 205		Glu	Thr	
Lys	Ile 210		Lys	Lys	Trp	Ala 215		Asp	Pro	Leu			Tyr	His	Pro	
Tyr 225		Phe	Gln	Val	Met 230		Asn	Val	Gly		220 Arg	Tyr	Arg	Phe		
	Tyr	Lys	Asn		Ala	Ile	Glu	Leu		235 Ala	Arg	Ile	Pro		240 Leu	
Thr	Asn	Asp	Tyr 260	245 Phe	Lys	Thr	Pro	Leu 265	250 Tyr	Thr	Leu	His	Phe 270	255 Lys	Arg	

Asn Ile Ser Val Tyr Leu Thr Ser Thr Tyr Asp Phe 275 280

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					gtt Val											206
					acg Thr 50											254
					aaa Lys											302
					gag Glu											350
					att Ile											398
					acc Thr											446
cat His 125	att Ile	ttc Phe	cct Pro	ttg Leu	atc Ile 130	gcc Ala	aaa Lys	gac Asp	ggg ggg	ggc Gly 135	gtg Val	tta Leu	gcg Ala	cgc Arg	acg Thr 140	494
ggc Gly	cat His	act Thr	gaa Glu	gcg Ala 145	agc Ser	gtg Val	gat Asp	ttg Leu	tgc Cys 150	aaa Lys	tta Leu	gct Ala	gga Gly	tta Leu 155	aag Lys	542

ccc gtg agc Pro Val Ser								590
aga agg ggg Arg Arg Gly 175	-		Asp Phe	_				638
aaa act ctc Lys Thr Leu 190	_		_	Tyr A	-	-	-	686
agt ttg ctg Ser Leu Leu 205		e Cys Gln						734
caa acg caa Gln Thr Gln	_		_					782
gct ttt aag Ala Phe Lys								830
cgt ttc cac Arg Phe His 255						_		878
ttt gaa gtg Phe Glu Val 270				Leu L				926
tat ttg atc Tyr Leu Ile 285		n Thr His		-				974
gat ttt ggg Asp Phe Gly								1022
ttc agg ctc Phe Arg Leu					yr Lys			1070
ggg ttt ggg Gly Phe Gly 335						aggctc		1116
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Ala Lys Lys Phe Glu Leu Pro Pro Met Val Ser Val Asn Asp Ser Asn
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His Glu Thr Ala Phe Thr Val Ser Ile Asp Ala Lys Glu Ala Arg Thr
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Gly Ile Ser Ala Phe Glu Arg His Leu Thr Ile Glu Leu Leu Cys Lys
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Asp Thr Thr Lys Pro Ser Asp Phe Val Arg Pro Gly His Ile Phe Pro
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Leu Ile Ala Lys Asp Gly Gly Val Leu Ala Arg Thr Gly His Thr Glu
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Ala Ser Val Asp Leu Cys Lys Leu Ala Gly Leu Lys Pro Val Ser Val
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Ile Cys Glu Ile Met Lys Glu Asp Gly Ser Met Ala Arg Arg Gly Asp
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Lys Phe Leu Ser Asp Phe Ala Leu Lys His Asn Leu Lys Thr Leu Tyr
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Val Ser Asp Leu Ile Ser Tyr Arg Leu Glu Asn Glu Ser Leu Leu Lys
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                        215
Tyr Thr Phe Leu Asp His Gln Gln Lys Asn His Tyr Ala Phe Lys Phe
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Lys Gly Ala Lys Thr His Asp Leu Ala Pro Leu Val Arg Phe His Pro
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                                   250
Ile Lys Glu Asp Phe Asp Phe Leu Thr Thr Asp Ala Phe Glu Val Phe
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Phe Lys Ala Leu Glu Tyr Leu Lys His Glu Gly Gly Tyr Leu Ile Phe
                           280
Met Asn Thr His Ser Lys Glu Asn Asn Val Val Lys Asp Phe Gly Ile
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                                           300
Gly Ala Leu Val Leu Lys Asn Leu Gly Ile Lys Asp Phe Arg Leu Leu
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											gaa Glu 30					152
_					-						att Ile					200
				_		_		-		-	agt Ser					248
											gtc Val					296
_				-				-			att Ile					341
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Phe																

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                        Gly Glu Asn Ile Leu Phe His Lys Leu Ile
tta aca tgc ttt tta geg ett gta gea ata acc att caa get tgc ggt
                                                                   159
Leu Thr Cys Phe Leu Ala Leu Val Ala Ile Thr Ile Gln Ala Cys Gly
                 15
                                     20
tat aaa gcc cct cca ttc aat gaa aaa ccc gct aaa aaa act tca aac
                                                                   207
Tyr Lys Ala Pro Pro Phe Asn Glu Lys Pro Ala Lys Lys Thr Ser Asn
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age tet aat tet tet atg caa acg eec ace aac age ace acg eea gaa
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Ser Ser Asn Ser Ser Met Gln Thr Pro Thr Asn Ser Thr Thr Pro Glu
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                             50
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Phe Leu Asn Gln Pro
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Phe Leu Asn Asn Lys His Arg Glu Lys Gly Leu Lys Ala Glu Glu Glu
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Ala Cys Gly Phe Leu Lys Ser Leu Gly Phe Glu Met Val Glu Arg Asn
                         25
     20
                                                                   200
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Phe Phe Ser Gln Phe Gly Glu Ile Asp Ile Ile Ala Leu Lys Lys Gly
                     40
                                          45
                                                                   248
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Val Leu His Phe Ile Glu Val Lys Ser Gly Glu Asn Phe Asp Pro Ile
tat gcg atc acg ccg agc aaa tta aaa aag atg att aaa acg atc cgc
                                                                   296
Tyr Ala Ile Thr Pro Ser Lys Leu Lys Lys Met Ile Lys Thr Ile Arg
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                                  75
tgt tat ttg tcc caa aaa gat ccc aat agc gat ttt tgc ata gac gct
                                                                   344
Cys Tyr Leu Ser Gln Lys Asp Pro Asn Ser Asp Phe Cys Ile Asp Ala
         85
                             90
                                                  95
ctt att gtg aaa aat ggt aaa ttt gag ctt tta gaa aat atc act ttt
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Leu Ile Val Lys Asn Gly Lys Phe Glu Leu Leu Glu Asn Ile Thr Phe
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                        105
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Lys Gly Val Leu His Phe Ile Glu Val Lys Ser Gly Glu Asn Phe Asp
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Pro Ile Tyr Ala Ile Thr Pro Ser Lys Leu Lys Lys Met Ile Lys Thr
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Ile Arg Cys Tyr Leu Ser Gln Lys Asp Pro Asn Ser Asp Phe Cys Ile

Asp Ala Leu Ile Val Lys Asn Gly Lys Phe Glu Leu Leu Glu Asn Ile 100 105 110

Thr Phe

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					agg Arg											592
					gac Asp 175											640
					gta Val											688
					gtt Val		-					-				736
					gat Asp				taag	gcaat	taa (catto	cttg	tt		783
tgg	cttta	aat a	attgi	tttt	ct aa	aaact	ttg	t tt	catg	gtaa	agct	tttta	ā			831
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Leu	Ile	Ser 35	Leu	Gly	Ala	Ser	Arg 40	Lys	Met	Gln	Met	Val 45	Pro	Met	Gly	
Leu	Gln 50		Val	Tyr	Glu	Ser 55	-	Ile	Ser	Ala,	Ile 60		Ser	Val	Ala	
Lys 65	Asp	Ile	Ile	Gly	Glu 70		Leu	Ala	Arg	Lys 75		Phe	Pro	Leu	Ala 80	
Gly	Thr	Ile	Ala	Leu 85	Tyr	Val	Phe	Phe	Ser 90		Met	Ile	Gly	Ile 95		
Pro	Gly	Phe	Glu 100		Pro	Thr	Ala	Ser 105		Ser	Phe	Thr	Leu 110		Leu	
Ala	Leu	Ile 115		Phe	Phe	Tyr	Tyr 120		Phe	Glu	Gly	Ile 125		Val	Gln	
Gly	Phe 130		Lys	Tyr	Phe	Ala 135		Phe	Ala	Gly	Pro 140		Lys	Trp	Leu	
Ala 145		Phe	Met	Phe	Pro 150		Glu	Ile	Ile	Ser 155		Phe	Ser	Arg	Ile 160	
	Ser	Leu	Ser	Phe 165	Arg	Leu	Phe	Gly	Asn 170		Lys	Gly	Asp	Asp 175		
Phe	Leu	Leu	Ile		Leu	Leu	Leu	Val		Tro	Ala	Val	Pro		Ala	

Pro Phe Met Val I 195 Met Ile Leu Thr T 210 Gly His 225	200)	205	
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gtt tta ttg ctc g Val Leu Leu Leu V 20				
tta gtg cgc ttg t Leu Val Arg Leu I 35				
agt aag ggt gcg a Ser Lys Gly Ala I				
agc cac cat aaa g Ser His His Lys V 70				
cac caa aac gcc t His Gln Asn Ala I 85		n His Ile Ala		=
gat ttg ctc atc t Asp Leu Leu Ile F 100				
gct cac gct tta g Ala His Ala Leu A 115				

gct tcc cct aaa atc cta gcc cct agc atg aac act aac atg ctc aat Ala Ser Pro Lys Ile Leu Ala Pro Ser Met Asn Thr Asn Met Leu Asn 135 140 145	488
tcc cct atc act caa agt aat tta aaa cgc ttg aaa gat tcc aac cac Ser Pro Ile Thr Gln Ser Asn Leu Lys Arg Leu Lys Asp Ser Asn His 150 155 160	536
att att tta gac acc aaa aac gcc ctt tta gca tgc gac act aaa ggc Ile Ile Leu Asp Thr Lys Asn Ala Leu Leu Ala Cys Asp Thr Lys Gly 165 170 175	584
gat ggg gcg atg gct gag cct tta gaa atc ctt ttt aaa gcc gct caa Asp Gly Ala Met Ala Glu Pro Leu Glu Ile Leu Phe Lys Ala Ala Gln 180 185 190	632
acg ctc cta aaa gac gct tat ttt gaa aac aga gaa gtc ata gtc atg Thr Leu Leu Lys Asp Ala Tyr Phe Glu Asn Arg Glu Val Ile Val Met 195 200 205 210	680
ggc ggc gcg agt ata gaa aag att gac agc gtt cga acg att agc aat Gly Gly Ala Ser Ile Glu Lys Ile Asp Ser Val Arg Thr Ile Ser Asn 215 220 225	728
act ttc tagcgggatt caagcgagcg ctttagcttt ggcgttatat tttaagggag Thr Phe	784
cca	
	787
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Lys Gly Asp Gly Ala Met Ala Glu Pro Leu Glu Ile Leu Phe Lys Ala
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                                                     190
            180
Ala Gln Thr Leu Leu Lys Asp Ala Tyr Phe Glu Asn Arg Glu Val Ile
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            Met Lys Pro Gln Asp Ile Glu Ile Val Gln Ser Val Leu
gag att aca gga ccg att aag cct act gaa gtg tat gat aaa gcc aaa
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Glu Ile Thr Gly Pro Ile Lys Pro Thr Glu Val Tyr Asp Lys Ala Lys
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gag ctt ttt gaa aaa ggt gag att aca aac atg ttt gat tgt ggg ggc
Glu Leu Phe Glu Lys Gly Glu Ile Thr Asn Met Phe Asp Cys Gly Gly
 30
                     35
                                          40
aaa acc ccg cac cag agc gtt agt tct tat att tat aca gcc tta aac
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Lys Thr Pro His Gln Ser Val Ser Ser Tyr Ile Tyr Thr Ala Leu Asn
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                                      55
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Lys Gly Glu Glu Leu Pro Phe Lys Lys Val Gln Glu Asn Pro Thr Leu
             65
atc gct tta aaa gac gcg gct aaa gag cta ggt tta gac gct caa aaa
                                                                   349
Ile Ala Leu Lys Asp Ala Ala Lys Glu Leu Gly Leu Asp Ala Gln Lys
         80
ata agc gct cca agc tct aaa atc gcg cat gaa agg gat ttg cac ccc
                                                                   397
Ile Ser Ala Pro Ser Ser Lys Ile Ala His Glu Arg Asp Leu His Pro
     95
                        100
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Phe Leu Thr Tyr Met Ala Ile Asn Asn Glu Asn Leu Lys Cys Tyr Thr
                    115
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													ggc Gly			493
													cac His 155			541
			_				_						gac Asp			589
			-		-		_	_			_		agc Ser			637
	-			_				-				_	tcg Ser		_	685
	_						_			-	_		aat Asn			733
													att Ile 235			781
													ttg Leu			829
													ctt Leu			877
													gat Asp			925
					-			_	_		_		gtt Val		_	973
					cct Pro							taaa	aato	gag		1019
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Glu Lys Gly Glu Ile Thr Asn Met Phe Asp Cys Gly Gly Lys Thr Pro
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His Gln Ser Val Ser Ser Tyr Ile Tyr Thr Ala Leu Asn Lys Gly Glu
Glu Leu Pro Phe Lys Lys Val Gln Glu Asn Pro Thr Leu Ile Ala Leu
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Lys Asp Ala Ala Lys Glu Leu Gly Leu Asp Ala Gln Lys Ile Ser Ala
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Pro Ser Ser Lys Ile Ala His Glu Arg Asp Leu His Pro Phe Leu Thr
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Tyr Met Ala Ile Asn Asn Glu Asn Leu Lys Cys Tyr Thr Lys Thr Ile
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Phe His Glu Glu Ser Ser Lys Ser Ile Lys Gly Met Asp Arg Trp Leu
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Tyr Pro Asp Met Val Gly Val Arg Phe Leu His Ala Glu Leu Ser Asn
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Glu Asn Leu Ile Ala Phe Ser Lys Lys Phe Asp Thr Leu Pro Ile Lys
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Leu Val Ser Phe Glu Leu Lys Lys Glu Ile Ser Val His Asn Cys Arg
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            180
Glu Cys Tyr Phe Gln Ala Ile Ser Asn Ser Ser Trp Ala Asn Glu Gly
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Tyr Leu Val Gly Arg His Ile Asp Thr His Asn Pro Gln Leu Met Asp
                        215
                                            220
Leu Leu Lys Arg Leu His Ala Ser Phe Gly Ile Gly Val Ile Asp Leu
                    230
                                        235
Arg Thr Asn Glu Asp Lys Ser Ala Ile Leu Leu Asn Ala Lys Tyr Lys
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                245
Glu Lys Ile Asp Tyr Thr Val Ala Ser Glu Leu Ser Ala Lys Asn Glu
                                265
Lys Phe Ser Gly Phe Leu Lys Ser Val Val Asp Tyr Asp Pro Asn His
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_					-		_	_	_	_	tac Tyr					309
											cat His					357
	ttt aaa gga cag tgc gca tgaaaaaaat cttagtcata ggcgatctga Phe Lys Gly Gln Cys Ala 85															405
tcg	ctgat	ta t	tatt	tgtg	gg gg	ggaag	gagc	g aad	egget	tt						444
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Arg	Ser	Tyr	Asn 20	Glu	Ile	Val	Ser	Ile 25	Leu	Lys	Glu	His	Leu 30	Gly	Asp	
Phe	Lys	Val 35		Tyr	Ile	Lys	Asn 40		Tyr	Ala	Phe	Phe 45		Lys	His	
Thr	Gln 50		His	Ile	Glu	Pro 55		Ile	Leu	Asp	Leu 60		Tyr	Thr	Pro	
Leu 65		Asp	Leu	Glu	Ser 70		Ile	Lys	Asp	Tyr 75	Leu	Pro	His	Ile	His 80	
	Ile	Phe	Lys	Gly 85	Gln	Cys	Ala			75					80	
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748

ttt aaa gag atg ata gat cta atc aaa taaggacaag aatgcgttat

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Phe Lys Glu Met Ile Asp Leu Ile Lys
185 190
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ttg aaa aac ctg gtt ttg tgc gat ata ggc aac acg cgt atc cat ttt 162 Leu Lys Asn Leu Val Leu Cys Asp Ile Gly Asn Thr Arg Ile His Phe

10			15					20					25	
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							att Ile							258
							aac Asn							306
		-		-		_	ggg							354
							gtg Val							402
							ggc Gly 115							450
							cat His							498
							tta Leu							546
							tac Tyr							594
							aat Asn							642
							ttt Phe 195							690
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Ser Ser Ala Lys Glu Asp Leu Lys Arg Leu Gly Ile Gln Lys Glu Ile
                            40
Phe Tyr Ile Ser Val Asn Glu Glu Asn Glu Lys Ala Leu Leu Asn Cys
                        55
Tyr Pro Asn Ala Lys Asn Ile Ala Gly Phe Phe His Leu Glu Thr Asp
Tyr Val Gly Leu Gly Ile Asp Arg Gln Met Ala Cys Leu Ala Val Asn
                                    90
                85
Asn Gly Val Val Val Asp Ala Gly Ser Ala Ile Thr Ile Asp Leu Ile
                                105
Lys Glu Gly Lys His Leu Gly Gly Cys Ile Leu Pro Gly Leu Ala Gln
                            120
                                                 125
Tyr Ile His Ala Tyr Lys Lys Ser Ala Lys Ile Leu Glu Gln Pro Phe
                        135
                                            140
Lys Ala Leu Asp Ser Leu Glu Val Leu Pro Lys Ser Thr Arg Asp Ala
                    150
                                        155
Val Asn Tyr Gly Met Val Leu Ser Val Ile Ala Cys Ile Gln His Leu
                                    170
                165
Ala Lys Asn Gln Lys Ile Tyr Leu Cys Gly Gly Asp Ala Lys Tyr Leu
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                                                     190
Ser Ala Phe Leu Pro His Ser Val Cys Lys Glu Arg Leu Val Phe Asp
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Gly Met Glu Ile Ala Leu Lys Lys Ala Gly Ile Leu Glu Cys Lys
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attatgtgat aatcaccaca agtaatcggc ttagtgtcat attacgaaga tttaagatca 180
taaaaggaaa aaag atg gtt aat aaa gat gtg aaa caa acc act gct ttt
                Met Val Asn Lys Asp Val Lys Gln Thr Thr Ala Phe
ggc gct ccc gtt tgg gat gac aac aat gtg att acg gcc ggc cct aga
                                                                   278
Gly Ala Pro Val Trp Asp Asp Asn Asn Val Ile Thr Ala Gly Pro Arg
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			gtg gtg cat gct Val Val His Ala 55	
			aaa gac atc act Lys Asp Ile Thr 70	
~ ~			aaa aaa acc gaa Lys Lys Thr Glu	-
=	Thr Val Ala		ggc agt gcg gat Gly Ser Ala Asp 105	
_		•	tac act gaa gaa Tyr Thr Glu Glu 120	
			ttc ttt atc cgt Phe Phe Ile Arg 135	

agc tta tac caa gta aca tgg gtt atg agc gat agg ggt att cct aaa 758 Ser Leu Tyr Gln Val Thr Trp Val Met Ser Asp Arg Gly Ile Pro Lys 175 tct ttc cgc cac atg gat ggt ttt ggc agc cac act ttc agt ctt atc 806 Ser Phe Arg His Met Asp Gly Phe Gly Ser His Thr Phe Ser Leu Ile 190 195 aac gcg aaa ggc gaa cgc ttt tgg gtg aaa ttc cac ttt cac acc atg 854 Asn Ala Lys Gly Glu Arg Phe Trp Val Lys Phe His Phe His Thr Met 210 215 caa ggc gtt aag cat ttg act aac gaa gaa gcc gca gaa gtt agg aag 902 Gln Gly Val Lys His Leu Thr Asn Glu Glu Ala Ala Glu Val Arg Lys 225 230

tat gat ccg gat tcc aat caa agg gat tta ttc aat gcg atc gct aga

aaa ttc cct gat ttc atc cac act caa aaa cga gat cct caa acc aat

Lys Phe Pro Asp Phe Ile His Thr Gln Lys Arg Asp Pro Gln Thr Asn

ttg cct aac cat gac atg gta tgg gat ttt tgg agt aat gtt cct gaa

Leu Pro Asn His Asp Met Val Trp Asp Phe Trp Ser Asn Val Pro Glu

165

150

145

160

662

710

Tyr	Asp	Pro	Asp 240	Ser	Asn	Gln	Arg	Asp 245	Leu	Phe	Asn	Ala	Ile 250	Ala	Arg	
					tgg Trp											998
~	_	_	_		cga Arg			_		-	_					1046
			_		cca Pro 290	_	_	_						_		1094
			_		tat Tyr		_	_		_			_		-	1142
_	_		-	-	cct Pro					_		_		_		1190
			_		tct Ser			_			_		_			1238
_					ata Ile	_	_				_	_				1286
					ggt Gly 370											1334
			_		agc Ser		_					_	_	_	_	1382
					ttc Phe											1430
					tac Tyr											1478
					cgc Arg											1526
					gag Glu 450											1574

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gag gga gtt aaa aaa gct ctt gaa aaa cac caa aaa atg atg aaa gac
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Glu Gly Val Lys Lys Ala Leu Glu Lys His Gln Lys Met Met Lys Asp
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                                485
atg cat gga aaa gac atg cac cac aca aaa aag aaa aag taaccctttt
                                                                   1719
Met His Gly Lys Asp Met His His Thr Lys Lys Lys
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Trp Asp Asp Asn Asn Val Ile Thr Ala Gly Pro Arg Gly Pro Val Leu
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                                25
Leu Gln Ser Thr Trp Phe Leu Glu Lys Leu Ala Ala Phe Asp Arg Glu
                            40
Arg Ile Pro Glu Arg Val Val His Ala Lys Gly Ser Gly Ala Tyr Gly
Thr Phe Thr Val Thr Lys Asp Ile Thr Lys Tyr Thr Lys Ala Lys Ile
Phe Ser Lys Val Gly Lys Lys Thr Glu Cys Phe Phe Arg Phe Ser Thr
                                    90
Val Ala Gly Glu Arg Gly Ser Ala Asp Ala Val Arg Asp Pro Arg Gly
                                105
Phe Ala Met Lys Tyr Tyr Thr Glu Glu Gly Asn Trp Asp Leu Val Gly
        115
                            120
Asn Asn Thr Pro Val Phe Phe Ile Arg Asp Ala Ile Lys Phe Pro Asp
                        135
                                            140
Phe Ile His Thr Gln Lys Arg Asp Pro Gln Thr Asn Leu Pro Asn His
                    150
                                        155
Asp Met Val Trp Asp Phe Trp Ser Asn Val Pro Glu Ser Leu Tyr Gln
                165
                                    170
Val Thr Trp Val Met Ser Asp Arg Gly Ile Pro Lys Ser Phe Arg His
                                185
Met Asp Gly Phe Gly Ser His Thr Phe Ser Leu Ile Asn Ala Lys Gly
        195
                            200
                                                205
Glu Arg Phe Trp Val Lys Phe His Phe His Thr Met Gln Gly Val Lys
                        215
His Leu Thr Asn Glu Glu Ala Ala Glu Val Arg Lys Tyr Asp Pro Asp
                    230
                                        235
Ser Asn Gln Arg Asp Leu Phe Asn Ala Ile Ala Arg Gly Asp Phe Pro
                245
                                    250
Lys Trp Lys Leu Ser Ile Gln Val Met Pro Glu Glu Asp Ala Lys Lys
```

265

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Tyr Arg Phe His Pro Phe Asp Val Thr Lys Ile Trp Tyr Leu Gln Asp
                            280
                                                 285
Tyr Pro Leu Met Glu Val Gly Ile Val Glu Leu Asn Lys Asn Pro Glu
                        295
Asn Tyr Phe Ala Glu Val Glu Gln Ala Ala Phe Ser Pro Ala Asn Val
                    310
                                        315
Val Pro Gly Ile Gly Tyr Ser Pro Asp Arg Met Leu Gln Gly Arg Leu
                325
                                    330
Phe Ser Tyr Gly Asp Thr His Arg Tyr Arg Leu Gly Val Asn Tyr Pro
                                345
Gln Ile Pro Val Asn Lys Pro Arg Cys Pro Phe His Ser Ser Ser Arg
                            360
                                                365
Asp Gly Tyr Met Gln Asn Gly Tyr Tyr Gly Ser Leu Gln Asn Tyr Thr
                        375
Pro Ser Ser Leu Pro Gly Tyr Lys Glu Asp Lys Ser Ala Arg Asp Pro
                    390
                                        395
Lys Phe Asn Leu Ala His Ile Glu Lys Glu Phe Glu Val Trp Asn Trp
                405
                                    410
Asp Tyr Arg Ala Asp Asp Ser Asp Tyr Tyr Thr Gln Pro Gly Asp Tyr
                                425
Tyr Arg Ser Leu Pro Ala Asp Glu Lys Glu Arg Leu His Asp Thr Ile
                            440
                                                445
Gly Glu Ser Leu Ala His Val Thr His Lys Glu Ile Val Asp Lys Gln
                        455
                                            460
Leu Glu His Phe Lys Lys Ala Asp Pro Lys Tyr Ala Glu Gly Val Lys
                    470
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Lys Ala Leu Glu Lys His Gln Lys Met Met Lys Asp Met His Gly Lys
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                                    490
Asp Met His His Thr Lys Lys Lys
            500
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aac ggc ttt gtg aat atc aat aac gaa aaa atg tct aaa agt ttg ggg
                                                                   104
Asn Gly Phe Val Asn Ile Asn Asn Glu Lys Met Ser Lys Ser Leu Gly
aat agc ttt ttt gtt aaa gac gct ctc aaa aac tat gat ggc gaa att
                                                                   152
Asn Ser Phe Phe Val Lys Asp Ala Leu Lys Asn Tyr Asp Gly Glu Ile
                         25
ttg cgc aat tac tta cta ggg gtg cat tat cgc tct gtt ttg aat ttc
                                                                   200
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Leu 35	Arg	Asn	Tyr	Leu	Leu 40	Gly	Val	His	Tyr	Arg 45	Ser	Val	Leu	Asn	Phe 50	
	_	-												atc Ile 65		248
-			_	_	_									cca Pro		296
														gtt Val		344
		_	_	_		_	_	_						gaa Glu		392
														tta Leu		440
														gac Asp 145		488
								_	_	_	_			gaa Glu		53.6
4												-		gat Asp		584
Leu														atc Ile		632
ttg Leu 195											aag Lys				٠	674
taaa	.cacc	tc c	aatt	ttac	c tt	ttta	caca	tto	ctago	caac	aact	ttca	agc a	att		727
<210 <211 <212 <213	> 20 > PF	8 RT	bact	er p	oylor	i									•	
<400 Met			Glv	Phe	Val	Asn	Ile	Asn	Asn	Glu	Lvs	Met	Ser	Lvs	Ser	
1 Leu				5					10					15		
Lou	0±3	******	20	- 11C	1116	vai	دلات	25 25	A10	ыeu	പുട	USII	30	veh	GTĀ	

```
Glu Ile Leu Arg Asn Tyr Leu Leu Gly Val His Tyr Arg Ser Val Leu
                            40
Asn Phe Asn Glu Glu Asp Leu Leu Val Ser Lys Lys Arg Leu Asp Lys
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Ile Tyr Arg Leu Lys Gln Arg Val Leu Gly Thr Leu Gly Gly Ile Asn
                    70
Pro Asn Phe Lys Lys Glu Ile Leu Glu Cys Met Gln Asp Asp Leu Asn
                                     90
Val Ser Lys Ala Leu Ser Val Leu Glu Ser Met Leu Ser Ser Thr Asn
                                105
Glu Lys Leu Asp Gln Asn Pro Lys Asn Lys Ala Leu Lys Gly Glu Ile
                            120
                                                 125
Leu Ala Asn Leu Lys Phe Ile Glu Glu Leu Leu Gly Ile Gly Phe Lys
                        135
Asp Pro Ser Ala Tyr Phe Gln Leu Gly Val Ser Glu Ser Glu Lys Gln
                    150
                                         155
Glu Ile Glu Asn Lys Ile Glu Glu Arg Lys Arg Ala Lys Glu Arg Lys
                                     170
                                                         1.75
                165
Asp Phe Leu Lys Ala Asp Ser Ile Arg Glu Glu Leu Leu Lys Gln Lys
                                185
                                                     190
Ile Ala Leu Met Asp Thr Pro Gln Gly Thr Ile Trp Glu Lys Phe Phe
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atg ctg acg att gaa acc agt aaa aaa ttt gat aag gat ctt aaa att
                                                                   107
Met Leu Thr Ile Glu Thr Ser Lys Lys Phe Asp Lys Asp Leu Lys Ile
                                     10
ctt gtt aaa aac ggg ttt gat tta aag ctt ttg tat aaa gtg gtt gga
                                                                   155
Leu Val Lys Asn Gly Phe Asp Leu Lys Leu Leu Tyr Lys Val Val Gly
             20
                                 25
aat tta gcc aca gag caa ccc cta gct ccc aaa tac aaa gac cac cca
                                                                   203
Asn Leu Ala Thr Glu Gln Pro Leu Ala Pro Lys Tyr Lys Asp His Pro
         35
                             40
ctc aaa ggc ggt tta aaa gat ttt agg gaa tgc cac tta aaa ccg gat
                                                                   251
Leu Lys Gly Gly Leu Lys Asp Phe Arg Glu Cys His Leu Lys Pro Asp
     50
                         55
tta ttg ctt gtc tat caa att aaa aaa caa gaa aac acc ctc ttt tta
                                                                   299
Leu Leu Val Tyr Gln Ile Lys Lys Gln Glu Asn Thr Leu Phe Leu
 65
                     70
```

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gta agg tta ggc agt cat agc gag ctg ttt tgaaccgccc acacccctta
                                                                   349
Val Arg Leu Gly Ser His Ser Glu Leu Phe
taacgcttaa accaactacc ccccttttt agggataaat ttagggttga aacaccgctt 409
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Leu Val Lys Asn Gly Phe Asp Leu Lys Leu Lyr Lys Val Val Gly
                                25
Asn Leu Ala Thr Glu Gln Pro Leu Ala Pro Lys Tyr Lys Asp His Pro
                                                 45
                            40
Leu Lys Gly Gly Leu Lys Asp Phe Arg Glu Cys His Leu Lys Pro Asp
Leu Leu Val Tyr Gln Ile Lys Lys Gln Glu Asn Thr Leu Phe Leu
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                                        75
Val Arg Leu Gly Ser His Ser Glu Leu Phe
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                   Val Leu Lys Leu Asn Leu Lys Lys Ser Phe Gln
                    1
                                                         10
aaa gat ttt gat aaa ttg ctt ttg aat ggg ttt gat gat agc gtt ttg
                                                                   158
Lys Asp Phe Asp Lys Leu Leu Leu Asn Gly Phe Asp Asp Ser Val Leu
             15
                                                     25
aat gaa gtc att cta acc tta aga aaa aaa gaa ccg cta gat cca caa
                                                                   206
Asn Glu Val Ile Leu Thr Leu Arg Lys Lys Glu Pro Leu Asp Pro Gln
         30
                             35
ttt caa gat cat gcc tta aag gga aag tgg aaa cct ttt agg gaa tgc
                                                                   254
Phe Gln Asp His Ala Leu Lys Gly Lys Trp Lys Pro Phe Arg Glu Cys
                         50
cac att aag cct gat gtt ttg ctt gtg tat tta gtg aaa gat gat gaa
                                                                   302
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His Ile Lys Pro Asp Val Leu Leu Val Tyr Leu Val Lys Asp Asp Glu 60 65 70 75 ctg att ttg tta agg tta ggc agt cat agc gag ctg ttt taatccaccc 351 Leu Ile Leu Leu Arg Leu Gly Ser His Ser Glu Leu Phe 80 acacccctta taacgcttaa accaaatcgc ttgcgctata atgaactgat attatatttt 411 425 aaaaggaata aaca <210> 82 <211> 88 <212> PRT <213> Helicobacter pylori <400> 82 Val Leu Lys Leu Asn Leu Lys Lys Ser Phe Gln Lys Asp Phe Asp Lys 10 Leu Leu Leu Asn Gly Phe Asp Asp Ser Val Leu Asn Glu Val Ile Leu 25 Thr Leu Arg Lys Lys Glu Pro Leu Asp Pro Gln Phe Gln Asp His Ala 40 Leu Lys Gly Lys Trp Lys Pro Phe Arg Glu Cys His Ile Lys Pro Asp 55 60 Val Leu Leu Val Tyr Leu Val Lys Asp Glu Leu Ile Leu Leu Arg 70 75 80 Leu Gly Ser His Ser Glu Leu Phe 85 <210> 83 <211> 844 <212> DNA <213> Helicobacter pylori <220> <221> CDS <222> (111) ... (779) <400> 83 gtcgttattc gcgcttaatg agaacagagg tttttaaaac tatggtttcg tttaggttta 60 attaaatttc gctacaatta aataaaaacg ataattttag agagattggc atg caa 116 Met Gln ggt tta tgg att tat cca gag gat aca gaa gtt tta ggg gtt gct tgt 164 Gly Leu Trp Ile Tyr Pro Glu Asp Thr Glu Val Leu Gly Val Ala Cys 5 10 aag agc ctt tta aaa gca cta acg cca cgc tat caa aaa gtc gcc ttg 212 Lys Ser Leu Leu Lys Ala Leu Thr Pro Arg Tyr Gln Lys Val Ala Leu 20 25 ttt tcg ccc att agt gga ggg tgt gag agc ttg gag gag tgc gag agc 260 Phe Ser Pro Ile Ser Gly Gly Cys Glu Ser Leu Glu Glu Cys Glu Ser

40

35

<400> 84

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Met Gln Gly Leu Trp Ile Tyr Pro Glu Asp Thr Glu Val Leu Gly Val

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Ala Cys Lys Ser Leu Leu Lys Ala Leu Thr Pro Arg Tyr Gln Lys Val
Ala Leu Phe Ser Pro Ile Ser Gly Gly Cys Glu Ser Leu Glu Glu Cys
                            40
Glu Ser Leu Asn Pro Leu Glu Phe His Ser Ala Ile Ser Lys Gln Lys
                        55
Ala Leu Glu Leu Ala Ser Thr Ala Gln Glu Glu Leu Leu Phe Glu Thr
                    70
                                        75
Ile Leu Lys Arg Tyr Asp Glu Leu Gln Ser Thr His Asp Phe Val Ile
                85
                                    90
Asn Leu Gly Cys Ala Pro Lys Phe Phe Leu Asn Ala Pro Leu Asp Leu
                                105
Asn Thr Ile Leu Ala Lys His Leu Asn Ala Ser Val Val Ala Val Ala
                            120
                                                 125
Gln Thr Ser Leu Glu Tyr Leu Lys Ala Met His Ser His Ile Leu Lys
                        135
Lys Glu Ala Pro Phe Ala Val Gly Leu Phe Ala Gly Glu Thr Leu Glù
                    150
                                        155
Lys Pro His Phe Leu Ser Met Ser Leu Cys Lys Gln Gln Cys Glu Leu
                165
                                    170
Glu Ala Asp Leu Ile Glu Ser Val Leu Gln Ile Lys Ser Glu Ile Ile
            180
                                185
Thr Pro Leu Ala Phe Gln Arg Gly Leu Glu Lys Lys Ala Lys Lys Gln
                            200
Ile Lys Lys Val Val Leu Pro Glu Ser Glu Lys Met Lys Gly Phe
                        215
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<211> 821
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<213> Helicobacter pylori
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<221> CDS
<222> (79)...(753)
<221> misc_feature
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agccagaaat tgggcaaa atg tgg aat ttt aac acc gaa tac agc agt cag
                    Met Trp Asn Phe Asn Thr Glu Tyr Ser Ser Gln
tat ttt gat ttt aga gcc gcc ggt ttt gtc caa ttg att tct aat tac
                                                                   159
Tyr Phe Asp Phe Arg Ala Ala Gly Phe Val Gln Leu Ile Ser Asn Tyr
             15
                                 20
atc aat caa ttt tct tca acg ctt ttt gta acc aac ttg ccc gca caa
                                                                   207
Ile Asn Gln Phe Ser Ser Thr Leu Phe Val Thr Asn Leu Pro Ala Gln
         30
                             35
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gat att att tat gtg cct ggt tat gaa gtt tca ggg acg gct aaa tac Asp Ile Ile Tyr Val Pro Gly Tyr Glu Val Ser Gly Thr Ala Lys Tyr 45 50 55	255
aag ggc ttt tct tta ggc ttg agc gtg gcg cga tca tgg cct tct tta Lys Gly Phe Ser Leu Gly Leu Ser Val Ala Arg Ser Trp Pro Ser Leu 60 65 70 75	303
aag ggg cgt ttg atc gct gat gtg tat gaa ttg gcg gcc acg aca ggc Lys Gly Arg Leu Ile Ala Asp Val Tyr Glu Leu Ala Ala Thr Thr Gly 80 85 90	351
aat gtg ttt att ttg acg gca agt tat aaa atc cca cgc act ggt ctt Asn Val Phe Ile Leu Thr Ala Ser Tyr Lys Ile Pro Arg Thr Gly Leu 95 100 105	399
agc atc act tgg ctt tca cgc ttc gtt acg gat ttg agt tat tgc tct Ser Ile Thr Trp Leu Ser Arg Phe Val Thr Asp Leu Ser Tyr Cys Ser 110 115 120	447
tat agc cct tat cgt aac ggc cct acg gat att gac aga cgg cct agt Tyr Ser Pro Tyr Arg Asn Gly Pro Thr Asp Ile Asp Arg Arg Pro Ser 125 130 135	495
aat tgc cct aaa acg ccc ggg att ttt cat gtt cat aaa ccc ggt tat Asn Cys Pro Lys Thr Pro Gly Ile Phe His Val His Lys Pro Gly Tyr 140 145 150 155	543
ggg gtg agc agt ttt ttt gta acc tac aaa ccc acc tat aag aag ctt Gly Val Ser Ser Phe Phe Val Thr Tyr Lys Pro Thr Tyr Lys Lys Leu 160 165 170	591
aaa ggg ttg agc ttg aat gcg gtg ttt aac aat gtt ttt aac caa caa Lys Gly Leu Ser Leu Asn Ala Val Phe Asn Asn Val Phe Asn Gln Gln 175 180 185	639
tat att gat caa gca agc ccg gtg atg agc cct gat gaa ccc aat caa Tyr Ile Asp Gln Ala Ser Pro Val Met Ser Pro Asp Glu Pro Asn Gln 190 195 200	687
gac aaa tac gca aga ggc atg gca gag cct ggc ttt aac gct aga ttt Asp Lys Tyr Ala Arg Gly Met Ala Glu Pro Gly Phe Asn Ala Arg Phe 205 210 215	735
gaa att tcc tat aag ttt taataatgga tctaaaaata aggatttcat Glu Ile Ser Tyr Lys Phe 220 225	783
gggtagcgga tctaatcaaa aataaaacat tctttaga	821
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Ser Thr Leu Phe Val Thr Asn Leu Pro Ala Gln Asp Ile Ile Tyr Val
                            40
Pro Gly Tyr Glu Val Ser Gly Thr Ala Lys Tyr Lys Gly Phe Ser Leu
                        55
                                             60
Gly Leu Ser Val Ala Arg Ser Trp Pro Ser Leu Lys Gly Arg Leu Ile
                    70
                                         75
Ala Asp Val Tyr Glu Leu Ala Ala Thr Thr Gly Asn Val Phe Ile Leu
                85
                                    90
Thr Ala Ser Tyr Lys Ile Pro Arg Thr Gly Leu Ser Ile Thr Trp Leu
            100
                                105
Ser Arg Phe Val Thr Asp Leu Ser Tyr Cys Ser Tyr Ser Pro Tyr Arg
Asn Gly Pro Thr Asp Ile Asp Arg Arg Pro Ser Asn Cys Pro Lys Thr
                        135
                                             140
Pro Gly Ile Phe His Val His Lys Pro Gly Tyr Gly Val Ser Ser Phe
                    150
                                        155
Phe Val Thr Tyr Lys Pro Thr Tyr Lys Lys Leu Lys Gly Leu Ser Leu
                165
                                    170
Asn Ala Val Phe Asn Asn Val Phe Asn Gln Gln Tyr Ile Asp Gln Ala
            180
                                185
Ser Pro Val Met Ser Pro Asp Glu Pro Asn Gln Asp Lys Tyr Ala Arg
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Gly Met Ala Glu Pro Gly Phe Asn Ala Arg Phe Glu Ile Ser Tyr Lys
                        215
                                             220
Phe
225
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                                                                   107
Met Asn Leu Asn Phe Met Pro Leu Leu His Ala Tyr Asn His Ala Ser
att gat ttt cat ttc aat tct agt gct agg gat ttt tgc gtg cat gaa
                                                                   155
Ile Asp Phe His Phe Asn Ser Ser Ala Arg Asp Phe Cys Val His Glu
             20
                                 25
gtg cct ttg tat gaa ttt agt aac acg ggc gaa cat gcc gtt att caa
                                                                   203
Val Pro Leu Tyr Glu Phe Ser Asn Thr Gly Glu His Ala Val Ile Gln
                             40
```

		ggt Gly												251
		gta Val												299
_		 ctg Leu 85	_											347
_		gaa Glu			_	-				_				395
	_	ttg Leu									_			443
-		cgc Arg			_	_				_				491
		aca Thr												539
_		ttt Phe 165		_		_			_			_		587
		 tta Leu							-			_		635
		gct Ala					_			_		_		683
		agc Ser		_		_		-				-	-	731
		gaa Glu												779
		act Thr 245												827
		ggc Gly												875

260		265	270
		gag ggc gaa agg t Glu Gly Glu Arg I	
		gac ggc aaa aaa g Asp Gly Lys Lys A	_
		aaa gaa ttc cag c Lys Glu Phe Gln F 315	
Ser Ser His Ala		tct agg cgg ttt t Ser Arg Arg Phe I 330	
		gtg aaa gaa aaa g Val Lys Glu Lys A 345	
		agt tat gcg agc g Ser Tyr Ala Ser A	
		aat aat gac gaa t Asn Asn Asp Glu F 380	5 5
	gtgga tgtcattaga		tgcgt gatttttgct 1272 ttagg aatagctctt 1332 1350
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Ile Asp Phe His 1	Phe Asn Ser Ser	Ala Arg Asp Phe C	ys Val His Glu 30
	Glu Phe Ser Asn	Thr Gly Glu His A	

Gln Ile Leu Gly Val Arg Ile Ala Glu Leu Gly Tyr Ala Gly Leu Lys 65 70 75 80

Asp Lys Asn Ala Leu Thr Thr Gln Phe Ile Ser Leu Pro Lys Lys Tyr 85 90 95

Ala Pro Leu Leu Glu Lys Asn Thr Ser Asn Phe Gln Glu Lys Asn Leu 100 105 110

Val Arg Lys Ser Gly Leu Ser Thr Leu Glu Met Leu Gln Ile Phe Ser

40

55

Lys Ile Leu Ser Leu Asn Tyr His His Asn Lys Ile Lys Leu Gly His

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120
                                                 125
        115
Leu Lys Gly Asn Arg Phe Phe Met Arg Phe Lys Lys Met Thr Pro Leu
                        135
                                            140
Asn Ala Gln Lys Thr Lys Gln Val Leu Glu Gln Ile Ala Gln Phe Gly
                    150
                                        155
Met Pro Asn Tyr Phe Gly Ser Gln Arg Phe Gly Lys Phe Asn Asp Asn
                                    170
                165
His Gln Glu Gly Leu Lys Ile Leu Gln Asn Gln Thr Lys Phe Ala His
                                185
Gln Lys Leu Asn Ala Phe Leu Ile Ser Ser Tyr Gln Ser Tyr Leu Phe
                            200
                                                205
Asn Ala Leu Leu Ser Lys Arg Leu Glu Ile Ser Lys Ile Ile Ser Ala
                        215
                                             220
Phe Ser Val Lys Glu Asn Leu Glu Phe Phe Lys Gln Lys Asn Leu Ser
                    230
                                        235
Val Asp Ser Asp Thr Leu Lys Thr Leu Lys Asn Gln Ala His Pro Phe
                                    250
                245
Lys Ile Leu Glu Gly Asp Val Met Cys His Tyr Pro Tyr Gly Lys Phe
                                265
Phe Asp Ala Leu Glu Leu Glu Lys Glu Gly Glu Arg Phe Leu Lys Lys
        2.75
                            280
                                                 285
Glu Val Ala Pro Thr Gly Leu Leu Asp Gly Lys Lys Ala Leu Tyr Ala
                        295
                                            300
Lys Asn Leu Ser Leu Glu Ile Glu Lys Glu Phe Gln His Asn Leu Leu
                    310
                                        315
Ser Ser His Ala Lys Thr Leu Gly Ser Arg Arg Phe Phe Trp Val Phe
                325
                                    330
Val Glu Asn Val Thr Ser Gln Tyr Val Lys Glu Lys Ala Gln Phe Glu
                                345
Leu Gly Phe Tyr Leu Pro Lys Gly Ser Tyr Ala Ser Ala Leu Leu Lys
                            360
Glu Ile Lys His Glu Lys Gly Glu Asn Asn Asp Glu Phe
                        375
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ccagatccct taaaa atg aaa ctc ccg gtc gtt gag agc ttt ttt tcc tta 111
                 Met Lys Leu Pro Val Val Glu Ser Phe Phe Ser Leu
caa ggt gaa gga aaa agg ata ggc aag ccc agt ctt ttt ttg cgc tta
                                                                   159
Gln Gly Glu Gly Lys Arg Ile Gly Lys Pro Ser Leu Phe Leu Arg Leu
ggg ggg tgt aac ctt tca tgc aag ggc ttt aat tgt aaa acc tta ttg
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Gly	Gly 30	Cys	Asn	Leu	Ser	Cys 35	Lys	Gly	Phe	Asn	Cys 40	Lys	Thr	Leu	Leu	
					aca Thr 50											255
					tgg Trp	_						_		_		303
					tta Leu											351
					cca Pro	_	_									399
_	-		~ ~		ttt Phe		_						_	_		447
					ttt Phe 130											495
				_	gtc Val								_	_	_	543
_					aaa Lys	_								-		591
-					ttt Phe	-			_			-	_			639
					agc Ser		_									687
					cct Pro 210											735
					gcc Ala											783
					atc Ile											828

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attttatgct gagcgtatcg ctgacaaccc tttagggttt atccaacgct tggatctttt 948
gcctagtatt agcgggttcg ttcaaaaatt gcgcgagcat ggcggggaat tttttgaaat 1008
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cgagctgtgc aa
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Leu Ser Cys Lys Gly Phe Asn Cys Lys Thr Leu Leu Asn Asp Glu Ile
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Leu Thr Gly Cys Asp Ser Leu Tyr Ala Val His Pro Lys Phe Lys Thr
                        55
Ser Trp Asp Tyr Tyr Asn Glu Pro Lys Pro Leu Ile Glu Arg Leu Glu
                    70
                                        75
Asp Leu Ala Pro Asn Tyr Lys Asp Phe Asp Phe Ile Leu Thr Gly Gly
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                                    90
Glu Pro Ser Leu Tyr Phe Asn Asn Pro Ile Leu Ile Ser Val Leu Glu
                                105
                                                     110
His Phe Tyr Arg Gln Lys Ile Pro Leu Cys Val Glu Ser Asn Gly Ser
                            120
                                                 125
Ile Phe Phe Glu Phe Ser Pro Ile Leu Lys Glu Leu His Phe Thr Leu
                        135
Ser Val Lys Leu Ser Phe Ser Leu Glu Glu Glu Ser Lys Arg Ile His
145
                    150
                                        155
Leu Lys Ala Leu Gln Asn Ile Leu Asn Asn Ala Lys Ser Ala His Phe
                                    170
Lys Phe Val Leu Glu Ser Gln Asn Ala Ala Gln Ser Ile Ile Glu Ile
                                185
                                                     190
Gln Ser Leu Leu Lys Gln Leu Ser Leu Lys Asn Asn Glu Ile Phe Leu
                            200
                                                205
Met Pro Leu Gly Thr Asn Asn Glu Leu Asp Lys Asn Leu Lys Thr
                        215
                                            220
Leu Ala Pro Leu Ala Ile Lys His Gly Phe Arg Leu Ser Asp Arg Leu
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His Ile Arg Leu Trp Asp Asn Gln Lys Gly Phe
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<222> (120)...(1559)
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gat	ttta	ttc (ggtg	ggati	tg to	cagca	atca	a gc	ctca	ttgt	tcc	tatt	agc (gttai	tttta	119
_		-			aaa	_	-	_		_						167
Met 1	Val	vaı	Pne	Thr 5	Lys	Arg	vai	Ala	10	ser	ьeu	Pne	Val	15	ше	
				J					10					13		
tta	gtg	agc	gct	gtt	tta	atg	cat	tcg	tta	cac	ctt	tcc	caa	ctc	gta	215
Leu	Val	Ser	Ala	Val	Leu	Met	His	Ser	Leu	His	Leu	Ser	Gln	Leu	Val	
			20					25					30			
~~~	+ - +	a + +	+-+	ast	aaa	a t a	o a t	+ 00	~++		+ > 0	205	+ > 0	~~~	999	263
_					Lys				_							203
014	- 7 -	35	- 1 -		טעט		40	501	V 4 1	1110	- 7 -	45	- 7 -	014	110	
-	_				ttc							_				311
Glu		Gly	Leu	Asn	Phe		Leu	Ser	Asn	Leu	-	Val	Phe	Gly	Phe	
	50					55					60					
tta	atc	ttt	tta	aac	gtc	tta	agc	caa	ata	att	tta	aaa	tcc	aat	agc	359
					Val											
65					70					75					80	
				_	aaa		-							-		407
vai	GIII	ASII	Pne	85	Lys	гуѕ	Ата	ьуѕ	90	TYL	ser	гуѕ	ASII	95	гÀг	
				0,5					50					73		
act	ccc	gaa	ttt	atc	gcc	ttt	ttt	tca	ggt	atc	att	att	ttt	gta	gat	455
Thr	Pro	Glu	Phe	Ile	Ala	Phe	Phe	Ser	Gly	Ile	Ile	Ile	Phe	Val	Asp	
			100					105					110			
gat	tat	+++	aac	acc	cta	acc	ata	aaa	caa	atc	tca	aan	tct	tta.	220	503
					Leu											505
_	_	115					120	1				125	~			
					aca											551
Asp	130	нтѕ	ASII	ser	Thr	135	GIU	Arg	ьeu	Ата	140	ше	тте	Asp	ser	
	130					133					740					
act	tca	gcg	ccg	gtg	tgc	ttg	cta	gtc	ccc	att	tct	agt	tgg	ggg	gcg	599
	Ser	Ala	Pro	Val	Cys	Leu	Leu	Val	Pro		Ser	Ser	Trp	Gly		
145					150					155					160	
tat	att	atα	aaa	atc	atg	aat	aac	gac	agc	tca	CCC	tta	tta	aaa	gat	647
					Met											01,
			_	165				-	170					175	-	
					gtg											695
ser	Pile	Ser	180	ьеu	Val	GIII	ser	185	ser	ser	Asn	Tyr	19r	Ата	IIe	
								100					100			
					gtc											743
Phe	Ala		Ile	Ala	Val	Phe		Thr	Ile	Leu	Trp		Ile	Asn	Leu	
		195					200					205				

	_	_	_	_						aag Lys 220	-				791
_		_	-	_				_		ttg Leu	_	_			839
				_	_					tca Ser	_				887
							_		-	ttt Phe	_				935
		_		_		_				tta Leu	-		_		983
	_			-				_	_	gat Asp 300			_	_	1031
										tgg Trp					1079
		-		_	_					ttg Leu	-			_	1127
						 				atg Met					1175
										ggc Gly					1223
										atg Met 380					1271
										ggc Gly					1319
										cta Leu					1367
										caa Gln					1415

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1463
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Thr Ile Ala Met Leu Cys Ser Ala Val Ser Leu Gly Val Ala Ser Phe
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atg tat tcg cgt tcg ctc gct ctt tta atc ggt gtg gct ttg ctt gtg
                                                                  1511
Met Tyr Ser Arg Ser Leu Ala Leu Leu Ile Gly Val Ala Leu Leu Val
    450
                        455
ggg gtg ttt tat ctt tta aaa aaa ttt tat ggt gaa aat cta aaa act
                                                                   1559
Gly Val Phe Tyr Leu Leu Lys Lys Phe Tyr Gly Glu Asn Leu Lys Thr
                    470
                                        475
tgaatattga ttgaagaagc ttaaaaatcc cattttttaa aattaaaata aggttttatc 1619
gatccctatt tgactcaaaa agagtcttat tccattatca atcaattaaa aaaggttatt 1679
caaaaataac catacaatta taaaaatctt c
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Glu Lys Gly Leu Asn Phe Asn Leu Ser Asn Leu Tyr Val Phe Gly Phe
                        55
                                            60
Leu Ile Phe Leu Gly Val Leu Ser Gln Val Ile Leu Lys Ser Gly Ser
                                        75
Val Gln Asn Phe Val Lys Lys Ala Lys Lys Tyr Ser Lys Asn Ala Lys
                85
                                    90
Thr Pro Glu Phe Ile Ala Phe Phe Ser Gly Ile Ile Ile Phe Val Asp
                                105
            100
                                                    110
Asp Tyr Phe Asn Ala Leu Thr Val Gly Gln Ile Ser Lys Ser Leu Asn
        115
                            120
                                                125
Asp Ala His Asn Ser Thr Arg Glu Arg Leu Ala Tyr Ile Ile Asp Ser
                        135
                                            140
Thr Ser Ala Pro Val Cys Leu Leu Val Pro Ile Ser Ser Trp Gly Ala
                    150
                                        155
Tyr Ile Met Gly Ile Met Asn Asn Asp Ser Ser Pro Leu Leu Lys Asp
                165
                                    170
                                                        175
Ser Phe Ser Val Leu Val Gln Ser Leu Ser Ser Asn Tyr Tyr Ala Ile
                                185
Phe Ala Leu Ile Ala Val Phe Leu Thr Ile Leu Trp Gln Ile Asn Leu
                            200
                                                205
Pro Ser Met Arg Lys Tyr Gln Asn Ile Gly Val Lys Asp Phe Tyr Ser
                        215
                                            220
Glu Glu Glu Ser Ser Ser Lys Leu Ala Pro Leu Ser Leu Leu Pro
                   230
                                        235
Leu Ser Ile Leu Leu Ile Val Ser Ile Ser Ser Leu Leu Phe Tyr
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Thr Gly Val Ile Leu Lys Asn Thr Asp Ala Ser Phe Ser Leu Phe Tyr
                                 265
            260
Gly Gly Leu Phe Ser Leu Ile Val Thr Tyr Leu Leu Ala Tyr Lys Phe
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                                                 285
        275
Leu Glu Lys Gly Ser Phe Phe Lys Leu Met Leu Asp Gly Phe Lys Ser
                        295
Val Gly Pro Ala Ile Leu Val Leu Thr Leu Ala Trp Ala Ile Gly Pro
                    310
                                         315
Val Ile Arg Asp Asp Ala Gln Thr Gly Leu Tyr Leu Ala Asn Ile Ser
                325
                                    330
Lys Gly Phe Leu Asn Asn Gly Gly Gly Val Tyr Met Pro Leu Ile Phe
            340
                                345
Phe Leu Ile Ser Gly Phe Ile Ala Phe Ser Thr Gly Thr Ser Trp Gly
                            360
Ala Phe Ala Ile Met Leu Pro Ile Gly Ala Gly Met Ala Ser Glu Ser
                        375
                                             380
Asp Ile Ile Leu Ile Val Ser Ala Ile Leu Ser Gly Ala Val Tyr Gly
                                         395
                    390
Asp His Thr Ser Pro Ile Ser Asp Thr Thr Ile Leu Ser Ala Thr Gly
                                    410
                405
Ala Gly Cys Ser Val Gln Ser His Phe Ile Thr Gln Leu Pro Tyr Ala
                                425
                                                     430
            420
Thr Ile Ala Met Leu Cys Ser Ala Val Ser Leu Gly Val Ala Ser Phe
        435
                            440
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Met Tyr Ser Arg Ser Leu Ala Leu Leu Ile Gly Val Ala Leu Leu Val
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Gly Val Phe Tyr Leu Leu Lys Lys Phe Tyr Gly Glu Asn Leu Lys Thr
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cattaaggtg ttttaagtta a att tcc tta tct gtt aaa cat acg gat aat
                        Ile Ser Leu Ser Val Lys His Thr Asp Asn
                         1
gtt ata tct tta agg aaa gaa aat ggg gtt agg aca cta ata agt tta
                                                                   159
Val Ile Ser Leu Arg Lys Glu Asn Gly Val Arg Thr Leu Ile Ser Leu
                 15
                                     20
                                                          25
ggg att ttg tta agc gtt ttg agt ggc gat gat ctg aag ttg tat tca
                                                                   207
Gly Ile Leu Leu Ser Val Leu Ser Gly Asp Asp Leu Lys Leu Tyr Ser
             30
                                 35
                                                      40
aaa ctt tca gtc tat tcg gct gga agt ggg atg att ggg att gat att
                                                                   255
Lys Leu Ser Val Tyr Ser Ala Gly Ser Gly Met Ile Gly Ile Asp Ile
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50 55 45 gac aaa cgg aca ttt tat aag cga gcg ttc gct ttc acg atg aaa tcg 303 Asp Lys Arg Thr Phe Tyr Lys Arg Ala Phe Ala Phe Thr Met Lys Ser 65 70 ttg ttc ggt gaa aac ttg ctt ttg ttt gtc aaa tta aag cat tct gcg 351 Leu Phe Gly Glu Asn Leu Leu Phe Val Lys Leu Lys His Ser Ala 80 85 ttg acg agc aaa cac atg aaa ggg cct tta gaa aac cgc cat cac cat 399 Leu Thr Ser Lys His Met Lys Gly Pro Leu Glu Asn Arg His His His 100 447 tct ttc act aaa aat tat gaa aaa gcg gtt aat ggt tgt caa aag tat Ser Phe Thr Lys Asn Tyr Glu Lys Ala Val Asn Gly Cys Gln Lys Tyr 110 115 495 ttc cat att aaa ttg cct gaa ggc gct cct agc aac ttc aaa tca ggt Phe His Ile Lys Leu Pro Glu Gly Ala Pro Ser Asn Phe Lys Ser Gly 125 130 545 tca tac atg gcc act atg gtg gtg cgt ttt taaagcgtta tttggggtat Ser Tyr Met Ala Thr Met Val Val Arg Phe 145 tetttaatae eettategte tittaaaata eeatettita aaageacaaa titatiitti 605 agcccttttt taaatcttct taaa 629 <210> 94 <211> 148 <212> PRT <213> Helicobacter pylori Ile Ser Leu Ser Val Lys His Thr Asp Asn Val Ile Ser Leu Arg Lys 10 Glu Asn Gly Val Arg Thr Leu Ile Ser Leu Gly Ile Leu Leu Ser Val 25 Leu Ser Gly Asp Asp Leu Lys Leu Tyr Ser Lys Leu Ser Val Tyr Ser 40 Ala Gly Ser Gly Met Ile Gly Ile Asp Ile Asp Lys Arg Thr Phe Tyr Lys Arg Ala Phe Ala Phe Thr Met Lys Ser Leu Phe Gly Glu Asn Leu 65 Leu Leu Phe Val Lys Leu Lys His Ser Ala Leu Thr Ser Lys His Met 85 90 Lys Gly Pro Leu Glu Asn Arg His His His Ser Phe Thr Lys Asn Tyr 105 Glu Lys Ala Val Asn Gly Cys Gln Lys Tyr Phe His Ile Lys Leu Pro 120 125

-120-

140

Glu Gly Ala Pro Ser Asn Phe Lys Ser Gly Ser Tyr Met Ala Thr Met

135

130

145

Val Val Arg Phe

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cataagactt caaagatttt ttaaagtttt aacattg atg cgt tgc gtg gtg tat
                                         Met Arg Cys Val Val Tyr
tct atc gct aaa agt tcg cct tta gag tta gtg aaa atc tat caa aag
                                                                   163
Ser Ile Ala Lys Ser Ser Pro Leu Glu Leu Val Lys Ile Tyr Gln Lys
                                                      20
             10
caa tgc agg caa ttt gat tgc gag ctg gaa ttg gtg gat tta ttc cct
                                                                   211
Gln Cys Arg Gln Phe Asp Cys Glu Leu Glu Leu Val Asp Leu Phe Pro
                             30
                                                                   259
aaa aat acc gcc aac gct caa aaa gtt tct aaa aaa ctg gct caa aaa
Lys Asn Thr Ala Asn Ala Gln Lys Val Ser Lys Leu Ala Gln Lys
agc tac tct cta gct ttt gag ccg tat tta aac cct aag gca aaa aat
                                                                   307
Ser Tyr Ser Leu Ala Phe Glu Pro Tyr Leu Asn Pro Lys Ala Lys Asn
 55
                     60
atc gcc tta cac cct aaa gct caa agg ggc gat agc ttt gcg ttt agt
                                                                   355
Ile Ala Leu His Pro Lys Ala Gln Arg Gly Asp Ser Phe Ala Phe Ser
                 75
aaa atg tta gaa aat cat ctt aat att aat ttt ttt atc gct gga gcg
                                                                   403
Lys Met Leu Glu Asn His Leu Asn Ile Asn Phe Phe Ile Ala Gly Ala
             90
tat ggg ttt gaa gaa aat ttt tta aag gat tgt caa gct tgg agt ttg
                                                                   451
Tyr Gly Phe Glu Glu Asn Phe Leu Lys Asp Cys Gln Ala Trp Ser Leu
        105
age gag atg act ttt age cat gaa gtg get aaa att gte tta tge gag
                                                                   499
Ser Glu Met Thr Phe Ser His Glu Val Ala Lys Ile Val Leu Cys Glu
    120
                        125
caa atc tat agg gct tta agc att att ttt aag cat cca tac cat aaa
                                                                   547
Gln Ile Tyr Arg Ala Leu Ser Ile Ile Phe Lys His Pro Tyr His Lys
                    140
                                        145
taggaggtgc gcatgcgttt ttacattatc tttacatttt tgtttattgt gggttttggt 607
gtgtttgttt atagtattg
                                                                   626
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<213> Helicobacter pylori
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Leu Val Asp Leu Phe Pro Lys Asn Thr Ala Asn Ala Gln Lys Val Ser
                            40
                                                 45
Lys Lys Leu Ala Gln Lys Ser Tyr Ser Leu Ala Phe Glu Pro Tyr Leu
Asn Pro Lys Ala Lys Asn Ile Ala Leu His Pro Lys Ala Gln Arg Gly
                    70
                                         75
Asp Ser Phe Ala Phe Ser Lys Met Leu Glu Asn His Leu Asn Ile Asn
Phe Phe Ile Ala Gly Ala Tyr Gly Phe Glu Glu Asn Phe Leu Lys Asp
            100
                                105
Cys Gln Ala Trp Ser Leu Ser Glu Met Thr Phe Ser His Glu Val Ala
                            120
                                                 125
Lys Ile Val Leu Cys Glu Gln Ile Tyr Arg Ala Leu Ser Ile Ile Phe
                        135
                                             140
Lys His Pro Tyr His Lys
145
                    150
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<222> (110)...(976)
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agcaaaactt ggatacaatc ctaacaaaac tgcaaaatta aggaaaaac atg gga ttt 118
                                                       Met Gly Phe
gca gat ttc ttt aaa aat ttt aag atc aat aaa ttg cgg aca gcg cca
                                                                   166
Ala Asp Phe Phe Lys Asn Phe Lys Ile Asn Lys Leu Arg Thr Ala Pro
                         10
agt aag gaa gaa cag cca agc cat tgg gtg aaa tgc cct aaa tgt tat
                                                                   214
Ser Lys Glu Glu Gln Pro Ser His Trp Val Lys Cys Pro Lys Cys Tyr
20
                     25
gcg tta atg tat cat aaa gaa gtg ttt agt aaa tac agc gtg tgt ttg
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Ala Leu Met Tyr His Lys Glu Val Phe Ser Lys Tyr Ser Val Cys Leu
                 40
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								gaa Glu					310
								aag Lys					358
								tat Tyr 95					406
		-			_		-	tca Ser			_		454
								gtg Val					502
_	_		 -					ggc Gly	-	_		_	550
								gcg Ala					598
	 -		 -	 _		-		act Thr 175		_		-	646
								ttg Leu					694
								tat Tyr			_	_	742
								gca Ala					790
								caa Gln					838
								tta Leu 255					886
								aag Lys					934

taaagtttta acattgatgc gttgcgtggt gtattctatc gctaaaagtt cgcctttaga 1036 gttagtgaaa atctatc 1053

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Phe

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gcttataata aacaaaatta gcttaagagt agtg atg caa ggg ttt ctt tta caa 115
                                      Met Gln Gly Phe Leu Leu Gln
aca caa agc ata aga gat gaa gat ttg atc gtg cgc gtt tta acc aaa
                                                                   163
Thr Gln Ser Ile Arg Asp Glu Asp Leu Ile Val Arg Val Leu Thr Lys
         10
                             15
aac cag ctc aaa acc ctc tat cgt ttc tat ggc aaa cgc cat agc gtg
                                                                   211
Asn Gln Leu Lys Thr Leu Tyr Arg Phe Tyr Gly Lys Arg His Ser Val
                         30
ctg aat gtg ggg cgt aaa att gat ttt gaa gaa gaa aac gat gat aag
                                                                   259
Leu Asn Val Gly Arg Lys Ile Asp Phe Glu Glu Glu Asn Asp Asp Lys
ttt tta ccc aag tta agg aat att ttg cat tta ggc tat att tgg gaa
                                                                   307
Phe Leu Pro Lys Leu Arg Asn Ile Leu His Leu Gly Tyr Ile Trp Glu
                 60
                                      65
aga gaa atg gag cgc ttg ttt ttt tgg caa cgc ttt tgc gct ctc ttg
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Arg Glu Met Glu Arg Leu Phe Phe Trp Gln Arg Phe Cys Ala Leu Leu
             75
                                 80
ttt agg cat tta gaa ggc gtg cat tct tta gat agc gtc tat ttt gac
                                                                   403
Phe Arg His Leu Glu Gly Val His Ser Leu Asp Ser Val Tyr Phe Asp
         90
                           • 95
act tta gat gat ggg gct aac aaa ctc gcc aaa cag cac ccc tta aga
                                                                   451
Thr Leu Asp Asp Gly Ala Asn Lys Leu Ala Lys Gln His Pro Leu Arg
    105
gtg att tta gaa atg tat gca acg ctt ttg aat ttt gaa ggg cgc ttg
                                                                   499
Val Ile Leu Glu Met Tyr Ala Thr Leu Leu Asn Phe Glu Gly Arg Leu
120
                    125
                                         130
caa agt tac aat tct tgt ttt tta tgc gat gca aaa tta gag cgt tct
                                                                   547
Gln Ser Tyr Asn Ser Cys Phe Leu Cys Asp Ala Lys Leu Glu Arg Ser
                                    145
gtc gct tta gcg caa ggg ttt att cta gcg cac ccc tct tgt ttg aaa
                                                                   595
Val Ala Leu Ala Gln Gly Phe Ile Leu Ala His Pro Ser Cys Leu Lys
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gct aaa agc cta aat tta gaa aaa atc caa gct ttt ttt cgc act caa

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Ala Lys Ser Leu Asn Leu Glu Lys Ile Gln Ala Phe Phe Arg Thr Gln
        170
                            175
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agc acg att gat tta gaa aca gaa gaa gta gaa gaa tta tgg cgc acg
Ser Thr Ile Asp Leu Glu Thr Glu Glu Val Glu Leu Trp Arg Thr
ctg aat tta ggg ttt tgaaaggtta aaaatgaaat ttaaattttt gaatatggat
                                                                   746
Leu Asn Leu Gly Phe
200
aatgaaagcg gttttatttt gattgaaaaa gaattgaaac gattaaacat tctcgctcaa 806
                                                                   810
gtca
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<213> Helicobacter pylori
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Ile Val Arg Val Leu Thr Lys Asn Gln Leu Lys Thr Leu Tyr Arg Phe
                                25
Tyr Gly Lys Arg His Ser Val Leu Asn Val Gly Arg Lys Ile Asp Phe
                            40
                                                 45
Glu Glu Glu Asn Asp Asp Lys Phe Leu Pro Lys Leu Arg Asn Ile Leu
                        55
His Leu Gly Tyr Ile Trp Glu Arg Glu Met Glu Arg Leu Phe Phe Trp
                    70
Gln Arg Phe Cys Ala Leu Leu Phe Arg His Leu Glu Gly Val His Ser
                                    90
                85
Leu Asp Ser Val Tyr Phe Asp Thr Leu Asp Asp Gly Ala Asn Lys Leu
                                105
Ala Lys Gln His Pro Leu Arg Val Ile Leu Glu Met Tyr Ala Thr Leu
                            120
                                                125
Leu Asn Phe Glu Gly Arg Leu Gln Ser Tyr Asn Ser Cys Phe Leu Cys
                        135
                                            140
Asp Ala Lys Leu Glu Arg Ser Val Ala Leu Ala Gln Gly Phe Ile Leu
                    150
                                        155
Ala His Pro Ser Cys Leu Lys Ala Lys Ser Leu Asn Leu Glu Lys Ile
                165
                                    170
Gln Ala Phe Phe Arg Thr Gln Ser Thr Ile Asp Leu Glu Thr Glu Glu
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Val Glu Glu Leu Trp Arg Thr Leu Asn Leu Gly Phe
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<213> Helicobacter pylori
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			_		-			_		-					ggt Gly		159
															aga Arg		207
				_		_	-				_		_	_	ttt Phe		255
															agg Arg 75		303
															acg Thr		351
										_					gta Val		399
															atc Ile		447
S													_		tat Tyr	_	495
_		_				_	_					_			agg Arg 155		543
															aaa Lys		591
															ttg Leu		639

tat ccc agc caa ttg att gaa gcg ttt tta gag ggg gtt atc gtg ttt 687

Tyr Pro Ser Gln Leu Ile Glu Ala Phe Leu Glu Gly Val Ile Val Phe

185

180

195

175

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tta atq qta atq tqq qct aaa aaa cac acc aaa acg cat ggg ttg ctg
                                                                   735
Leu Met Val Met Trp Ala Lys Lys His Thr Lys Thr His Gly Leu Leu
205
                    210
                                         215
att gtg gtt tat ggt ttg ggg tat tcc ttg atg cgc ttt att gcg gaa
                                                                   783
Ile Val Val Tyr Gly Leu Gly Tyr Ser Leu Met Arg Phe Ile Ala Glu
                225
ttt tac aga gag ccg gac agc caa atg ggg gtt tat ttt tta aat ttg
                                                                   831
Phe Tyr Arg Glu Pro Asp Ser Gln Met Gly Val Tyr Phe Leu Asn Leu
            240
                                 245
agc atg ggg cag att tta agc tta ttt atg gta att gtt tcg tta ggg
                                                                   879
Ser Met Gly Gln Ile Leu Ser Leu Phe Met Val Ile Val Ser Leu Gly
        255
                            260
                                                 265
                                                                   927
att tta ttg tat get aca aaa aat tet aaa aaa ata aag gaa aat caa
Ile Leu Leu Tyr Ala Thr Lys Asn Ser Lys Lys Ile Lys Glu Asn Gln
                        275
                                             280
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tgtttgatag cc
                                                                   999
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Ser Leu Gly Ser Ile Glu Val His Trp Tyr Gly Leu Ala Tyr Ala Cys
Ala Ile Val Thr Ala Phe Tyr Met Ala Leu Arg Met Ile Gln Lys Asp
                            40
Pro Lys Arg Phe Pro Ile Glu Arg Lys Glu Phe Glu Ser Tyr Phe Leu
                                             60
Trp Ala Glu Leu Gly Ile Val Leu Gly Ala Arg Ile Gly Tyr Ile Leu
                    70
                                        75
Ile Tyr Glu Pro Asn Ser Gly Tyr Tyr Leu Thr His Phe Trp Gln Ile
                                    90
Phe Asn Pro Phe Asp Ser His Gly Asn Phe Val Gly Ile Arg Gly Met
                                105
Ser Tyr His Gly Gly Leu Val Gly Phe Leu Ile Ala Ser Tyr Leu Tyr
                            120
                                                 125
Ser Arg Lys Asp Leu Lys Lys Leu Leu Ile Tyr Leu Asp Leu Ile Ala
                        135
                                             140
Ile Ser Leu Pro Leu Gly Tyr Val Phe Gly Arg Ile Gly Asn Phe Leu
                    150
                                        155
Asn Gln Glu Leu Val Gly Arg Ile Val Pro Lys Asp Ser His Leu Gly
                165
                                    170
Gln Ile Ile Gly Ile Met Val Asp Asn Glu Leu Arg Tyr Pro Ser Gln
                                185
Leu Ile Glu Ala Phe Leu Glu Gly Val Ile Val Phe Leu Met Val Met
```

															-	
Ψrn	Ala	195	Larg	ніс	ጥኮድ	Lare	200 Thr	Hic	Gly	T.e.u	T.eu	205 Tle	Val	Va1	ጥህድ	
	210					215					220					
Gly 225	Leu	Gly	Tyr	Ser	Leu 230	Met	Arg	Phe	Ile	A1a 235	GIu	Phe	Tyr	Arg	G1u 240	
Pro	Asp	Ser	Gln	Met 245	Gly	Val	Tyr	Phe	Leu 250	Asn	Leu	Ser	Met	Gly 255	Gln	
Ile	Leu	Ser	Leu 260		Met	Val	Ile	Val 265		Leu	Gly	Ile	Leu 270		Tyr	
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	2> (8		(9	94)												
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			,								Ser 5					
					•											1.50
Lys	cta Leu															160
10					15					20					25	
	gat Asp															208
-4-				30	0-1		<b>01</b>		35				****	40	200	
	agt															256
Arg	Ser	Leu	Asp 45	Lys	Lys	Pro	Trp	Asn 50	Val	Ala	Tyr	Val	Ala 55	Pro	Ser	
aga	agg	cct	act	gat	ggg	cgc	tat	ggg	gaa	aac	cct	aac	cgc	ttg	ggg	304
	Arg															
+	<b></b>															252
	tat Tyr					Val					Ser					352
	75					80					85					
	gaa Glu															400
90		-	-		95				<b>-</b>	100	<b>1</b>	_ •			105	
	cat															448
GIU	His	ASD	тте	110	rne	val	GIU	ASP	115	Trp	GIU	ser	rro	Thr 120	ьeu	

ggg gca	tgg	ggg	ctt	ggc	tgg	gaa	gtg	tgg	ctt	gat	ggc	atg	gaa	gtt	496
Gly Ala	Trp	Gly 125	Leu	Gly	Trp	Glu	Val 130	Trp	Leu	Asp	Gly	Met 135	Glu	Val	
acg caa Thr Glr															544
att cct Ile Pro 155	Val														592
caa aaa Gln Lys 170															640
gac ago Asp Ser															698
agc aag Ser Lys															736
ttt aaa Phe Lys		-		-	_	_	_		_	_	_		_		784
ccc ttg Pro Leu 235	Pro														832
att tta Ile Leu 250															880
att tta Ile Leu			Arg		Leu	Ala	Lys	Gly	Cys	Ala	Leu	Leu	Tyr	Lys	928
gaa caa Glu Glr															976
gct gaa Ala Glu					taag	ggaaq	gtg t	tggt	agtt	it to	gaato	egeet	=		1024
ttcgcct tagtgaa					atcat	ggga	a taa	atago	ggg	ttga	aatgt	gg g	ggagt	gaaaa	1084 1104
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## <213> Helicobacter pylori

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Gly Thr Phe His Pro Ala Thr Leu Leu Arg Ser Leu Asp Lys Lys Pro
                            40
Trp Asn Val Ala Tyr Val Ala Pro Ser Arg Arg Pro Thr Asp Gly Arg
                        55
                                            60
Tyr Gly Glu Asn Pro Asn Arg Leu Gly Ser Tyr Tyr Gln Phe Gln Val
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                                        75
Val Ile Lys Pro Ser Pro Ser Asn Ile Gln Glu Leu Tyr Leu Lys Ser
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Leu Glu Val Leu Gly Ile Asn Leu Asn Glu His Asp Ile Arg Phe Val
            100
                                105
Glu Asp Asn Trp Glu Ser Pro Thr Leu Gly Ala Trp Gly Leu Gly Trp
                            120
                                                125
Glu Val Trp Leu Asp Gly Met Glu Val Thr Gln Phe Thr Tyr Phe Gln
                        135
                                            140
Gln Val Gly Gly Ile Ala Cys Ser Pro Ile Pro Val Glu Ile Thr Tyr
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Gly Leu Glu Arg Leu Ala Met Tyr Val Gln Lys Val Glu Asn Ile Leu
                165
                                    170
Glu Ile Glu Trp Ala Lys Lys Asn His Asp Ser Val Asn Tyr Ala Gln
                                185
Val His Leu Glu Ser Glu Tyr Glu Phe Ser Lys Tyr His Phe Glu Thr
        195
                            200
                                                205
Ala Ser Val Lys Arg Leu Leu Glu Met Phe Lys Asn Ala Gln Ala Glu
                        215
                                            220
Ala Leu His Cys Leu Glu Asn Lys Leu Pro Leu Pro Ala Tyr Asp Phe
                    230
                                        235
Val Met Leu Cys Ser His Phe Phe Asn Ile Leu Asp Ala Arg Lys Ala
                                    250
                245
Ile Ser Val Ala Glu Arg Gln Asn Tyr Ile Leu Gln Ile Arg Asp Leu
            260
                                265
                                                    270
Ala Lys Gly Cys Ala Leu Leu Tyr Lys Glu Glu Glu Glu Glu Arg Glu
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Glu Arg Leu Lys Asn Ala Leu Thr Lys Ala Glu Asn Gly Val Ser
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_	et Al	la G	ln Ly	nr Le 5	eu Le	eu I	le I		nr As 10	sp G	ly I	le G	ly Ty 1	r .5
												ccc Pro 30		158
												acg Thr		206
 _	_	-			_			_				gaa Glu		254
	_	_		 _							_	tta Leu	-	302
			_			-	-					ccc Pro	-	350
												ggt Gly 110		398
												gct Ala		446
		_	-					-	-	-		tta Leu		494
												tta Leu		542
												ata Ile		590
												att Ile 190		638
												agc Ser		686
												gaa Glu		734

									atg Met 235					782
									gct Ala					830
 _	-					_	~ ~		aag Lys	_		_		878
			_		_	_			gat Asp					926
									aac Asn					974
 _				_			_		atc Ile 315	-	_		-	1022
		_							gga Gly			_		1070
	_							_	cct Pro		_			1118
									gta Val					1166
									att Iļe					1214
									gaa Glu 395					1262
									atc Ile					1310
			-	_				_	gat Asp				_	1358
									acc Thr					1406

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ggg age gtg tat tgc ttt gtt tta ggg gat gga gtc aaa tcc ata aaa
                                                                   1454
Gly Ser Val Tyr Cys Phe Val Leu Gly Asp Gly Val Lys Ser Ile Lys
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aac gga gcc tta aac aat atc gct agc agc gtg tta aaa ctc atg ggc
                                                                   1502
Asn Gly Ala Leu Asn Asn Ile Ala Ser Ser Val Leu Lys Leu Met Gly
    465
                        470
ctt aaa gcc cca gca acg atg gac gaa ccc cta ttt taaactaaag
                                                                   1548
Leu Lys Ala Pro Ala Thr Met Asp Glu Pro Leu Phe
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gaaaagaatg caaattgatg acgcattatt gcaacgcttg gaaaaattga gcatgctaga 1608
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gattaaagat ga
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                                25
Asp Leu Met Phe Lys Thr Leu Pro Tyr Ser Leu Ile Asp Thr His Gly
Leu Ser Val Gly Leu Pro Lys Gly Gln Met Gly Asn Ser Glu Val Gly
                        55
His Met Cys Ile Gly Ala Gly Arg Val Leu Tyr Gln Asp Leu Val Lys
                    70
                                        75
Ile Ser Leu Ser Leu Gln Asn Asp Glu Leu Lys Asn Asn Pro Ala Phe
                                    90
Leu Asn Thr Ile Gln Lys Ser Pro Val Val His Leu Met Gly Leu Met
                                105
                                                     110
Ser Asp Gly Gly Val His Ser His Ile Glu His Phe Ile Ala Leu Ala
                            120
        115
Leu Glu Cys Glu Lys Ser His Lys Lys Val Cys Leu His Leu Ile Thr
                        135
                                            140
Asp Gly Arg Asp Val Ala Pro Lys Ser Ala Leu Thr Tyr Leu Lys Gln
                    150
                                        155
Met Gln Asn Ile Cys Asn Glu Ser Ile Gln Ile Ala Thr Ile Ser Gly
                165
                                    170
Arg Phe Tyr Ala Met Asp Arg Asp Lys Arg Phe Glu Arg Ile Glu Leu
                                185
                                                     190
Ala Tyr His Ser Leu Met Gly Leu Asn His Thr Pro Leu Ser Pro Ser
                            200
                                                 205
Glu Tyr Ile Gln Ser Gln Tyr Asp Lys Asn Ile Thr Asp Glu Phe Ile
                        215
                                             220
Met Pro Ala Cys Phe Lys Asn Tyr Cys Gly Met Gln Asp Asp Glu Ser
                    230
                                        235
Phe Ile Phe Ile Asn Phe Arg Asn Asp Arg Ala Arg Glu Ile Val Ser
                245
                                    250
Ala Leu Gly Gln Lys Gln Phe Ser Gly Phe Lys Arg Gln Val Phe Lys
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Lys Leu His Ile Ala Thr Met Thr Pro Tyr Asp Asn Thr Phe Pro Tyr
                            280
Pro Val Leu Phe Pro Lys Glu Ser Val Gln Asn Thr Leu Ala Glu Val
                        295
                                             300
Val Ser Gln His Asn Leu Thr Gln Ser His Ile Ala Glu Thr Glu Lys
                    310
                                        315
Tyr Ala His Val Thr Phe Phe Ile Asn Gly Gly Val Glu Thr Pro Phe
                325
                                    330
                                                         335
Lys Asn Glu Asn Arg Val Leu Ile Gln Ser Pro Lys Val Thr Thr Tyr
            340
                                345
                                                     350
Asp Leu Lys Pro Glu Met Ser Ala Lys Glu Val Thr Leu Ala Val Leu
        355
                            360
                                                 365
Glu Gln Met Lys Leu Gly Thr Asp Leu Ile Ile Val Asn Phe Ala Asn
                        375
                                             380
Gly Asp Met Val Gly His Thr Gly Asn Phe Glu Ala Ser Val Lys Ala
                    390
                                         395
Val Glu Ala Val Asp Ala Cys Leu Gly Glu Ile Leu Ser Leu Ala Lys
                405
                                    410
Lys Leu Asp Tyr Ala Met Leu Leu Thr Ser Asp His Gly Asn Cys Glu
            420
                                425
Arg Met Lys Asp Glu Asn Gln Asn Pro Leu Thr Asn His Thr Ala Gly
                            440
        435
                                                 445
Ser Val Tyr Cys Phe Val Leu Gly Asp Gly Val Lys Ser Ile Lys Asn
                        455
                                            460
Gly Ala Leu Asn Asn Ile Ala Ser Ser Val Leu Lys Leu Met Gly Leu
                    470
                                         475
Lys Ala Pro Ala Thr Met Asp Glu Pro Leu Phe
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gagtttttaa c atg aat ttt caa gaa aat tta gcc gct ttg gat ttg gag
             Met Asn Phe Gln Glu Asn Leu Ala Ala Leu Asp Leu Glu
              1
tat ctt tgg cac cct tgt tcg caa atg caa gag cat caa aat ttc ccc
                                                                   158
Tyr Leu Trp His Pro Cys Ser Gln Met Gln Glu His Gln Asn Phe Pro
     15
                         20
att atc ccc att aaa aag gct caa ggg att tac ctc tat gat ttt aat
                                                                   206
Ile Ile Pro Ile Lys Lys Ala Gln Gly Ile Tyr Leu Tyr Asp Phe Asn
                     35
                                         40
gat aac gct tac atg gat ttg atc agc tca tgg tgg gtg aat ctt ttt
                                                                   254
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Asp	Asn	Ala	Tyr	Met 50	Asp	Leu	Ile	Ser	Ser 55	Trp	Trp	Val	Asn	Leu 60	Phe	
	cat His															302
-	tta Leu			_		_	_									350
	ctc Leu 95															398
	gcg Ala	-														446
	cac His	_				_			_	_	_		_			494
	tcg Ser															542
	ggc Gly	_						_						_		590
	aat Asn 175															638
_	ttg Leu		-		_	_	_		~	_		_	_	_		686
	gct Ala															734
	tat Tyr															782
	aac Asn															830
	ggg Gly 255															878

tta tgc ttg tct aag gg Leu Cys Leu Ser Lys G 270 2	y Ile Ser Gly		_
cta tta acc cat aat ga Leu Leu Thr His Asn G 290	u Ile Tyr Asn		
gaa aat aaa gcg ttt t Glu Asn Lys Ala Phe Lo 305			n Ala Leu
gca tgc gca tgc gcg a Ala Cys Ala Cys Ala A 320			
gtt att gaa aag aac a Val Ile Glu Lys Asn Ly 335			<del>-</del>
caa aac gca tta aaa c Gln Asn Ala Leu Lys P 350 3	o Leu Met Glu		
agg cat ttg ggc atg g Arg His Leu Gly Met Va 370	l Phe Ala Phe		
gag cgt ttg agt ttg gg Glu Arg Leu Ser Leu A 385			Gly Leu
tta tta cgc cct tta aa Leu Leu Arg Pro Leu Aa 400		_	
atc acg cat gaa gaa g Ile Thr His Glu Glu Va 415			
ctt gat gag tta aga ag Leu Asp Glu Leu Arg Ly 430 43	s Gly	tt tttaaaaata aaata	agtaaa 1409
aagcttgaat tggatcaagc	gatagctttt t		1440
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<400> 108 Met Asn Phe Gln Glu As	ח דא מום אום ז	Leu Jan Leu Clu Tre	Tou Tro
Met Asn Phe Gln Glu As 1 5 His Pro Cys Ser Gln Me	:	10	15
ITO CAS DEL GIN ME	c Jin Jiu nis (	OTH ASH THE FIG TIE	TIC LIO

			20					25					30		
Tle	Lys	Lvs		Gln	Glv	Ile	Tvr		Tvr	asp	Phe	Asn		Asn	Ala
	_	35					40					45			
Tyr	Met 50	Asp	Leu	Ile	Ser	Ser 55	Trp	Trp	Val	Asn	Leu 60	Phe	Gly	His	Asn
Asn 65	Ala	Tyr	Ile	Ser	Gln 70	Gln	Leu	Lys	Asn	Gln 75	Ile	Asp	Asp	Leu	Glu 80
His	Val	Leu	Leu	Ala 85	Ser	Phe	Ser	His	Lys 90	Pro	Ile	Ile	Thr	Leu 95	Ser
Gln	Arg	Leu	Cys 100	Gln	Leu	Thr	His	Met 105	Asp	Lys	Суѕ	Phe	Tyr 110	Ala	Asp
Asn	Gly	Ser 115	Ser	Суѕ	Val	Glu	Ile 120	Ala	Leu	Lys	Met	Ser 125	Tyr	His	Ala
His	Phe 130	Leu	Lys	Asn	Gln	Thr 135	Arg	Arg	Lys	Lys	Leu 140	Phe	Leu	Ser	Leu
Ser 145	Asn	Ser	Tyr	His	Gly 150	Glu	Thr	Leu	Gly	Ala 155	Leu	Ser	Val	Gly	Asp 160
Val	Lys	Leu	Tyr	Lys 165	Asp	Thr	Tyr	Thr	Pro 170	Leu	Leu	Leu	ГЛЗ	Asn 175	Leu
Thr	Thr	Pro	Val 180	Pro	Lys	Asn	Asp	His 185	Glu	Ile	Glu	Asn	Ser 190	Leu	Asn
Ala	Leu	Lys 195	Arg	Leu	Leu	Asp	Lys 200	His	Ser	Glu	Glu	Ile 205	Суѕ	Ala	Phe
Ile	Ala 210	Glu	Pro	Leu	Leu	Gln 215	Cys	Ala	Gly	Asn	Met 220	His	Ile	Tyr	Ser
Ala 225	Arg	Tyr	Leu	Lys	Gln 230	Ala	Val	Leu	Leu	Cys 235	Lys	Gln	Lys	Asn	Ile 240
His	Ile	Ile	Phe	Asp 245	Glu	Ile	Ala	Thr	Gly 250	Phe	Gly	Arg	Thr	Gly 255	Ser
Met	Phe	Ala	Tyr 260	Glu	Gln	Cys	Glu	Ile 265	Lys	Pro	Asp	Phe	Leu 270	Cys	Leu
Ser	Lys	Gly 275	Ile	Ser	Gly	Gly	Tyr 280	Leu	Pro	Leu	Ser	Ala 285	Leu	Leu	Thr
His	Asn 290	Glu	Ile	Tyr	Asn	Gln 295	Phe	Tyr	Ala	Pro	Tyr 300	Glu	Glu	Asn	Lys
Ala 305	Phe	Leu	His	Ser	His 310	Ser	Tyr	Thr	Gly	Asn 315	Ala	Leu	Ala	Cys	Ala 320
Cys	Ala	Asn	Ala	Thr 325	Leu	Asp	Ile	Phe	Glu 330	Lys	Glu	Asn	Val	Ile 335	Glu
Lys	Asn	Lys	Ala 340	Leu	Ser	Gly	Phe	Ile 345	Phe	Asn	Thr	Leu	Gln 350	Asn	Ala
Leu	Lys	Pro 355	Leu	Met	Glu	Gln	Gln 360	Val	Val	Ser	Asp	Leu 365	Arg	His	Leu
Gly	Met 370	Val	Phe	Ala	Phe	Glu 375	Val	Phe	Ile	Gln	Thr 380	Lys	Glu	Arg	Leu
Ser 385	Leu	Ala	Val	Phe	Lys 390	Lys	Thr	Leu	Lys	Lys 395	Gly	Leu	Leu	Leu	Arg 400
	Leu	Asn	Asn	Thr 405		Tyr	Leu	Met	Pro 410		Tyr	Ile	Ile	Thr 415	
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Leu	Arg	Lys 435	Gly												

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                        Met Gly Lys Met Lys Gln Glu Thr Ala Ile
gac tat gaa aaa tta gcg aat cat tgg aat aat aat gat gaa aac agc
                                                                   159
Asp Tyr Glu Lys Leu Ala Asn His Trp Asn Asn Asp Glu Asn Ser
                                     20
                 15
gaa gca cta aac gct ttt gca gac gct tac ctt tat aaa cat gag aaa
                                                                   207
Glu Ala Leu Asn Ala Phe Ala Asp Ala Tyr Leu Tyr Lys His Glu Lys
             30
                                 35
aag agt caa aag att cgg gca ata gag ata agt tct cta aac aaa gcc
                                                                   255
Lys Ser Gln Lys Ile Arg Ala Ile Glu Ile Ser Ser Leu Asn Lys Ala
         45
                             50
tgc atg gga gaa ttt tac cac aaa aac cca aaa tta ttt taataacgat
                                                                   304
Cys Met Gly Glu Phe Tyr His Lys Asn Pro Lys Leu Phe
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                         65
                                                                   360
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aga cat agt att tcg caa aga gct ttt gaa ttt ttt aag tat atg aat

Arg	His	Ser 195	Ile	Ser	Gln	Arg	Ala 200	Phe	Glu	Phe	Phe	Lys 205	Tyr	Met	Asn	
											atg Met 220					733
_				_	-	_			-		att Ile	_				781
_		_		_							aac Asn					829
		-			_						gag Glu				-	877
	_		_			_		-	_		cga Arg		-			925
		-		_	-				-		aaa Lys 300				•	973
											ggg Gly					1021
			_	_	-		_				ttg Leu	_		_		1069
						_		_	_	_	ata Ile	-		_		1117
											caa Gln					1165
											ttt Phe 380					1213
											atg Met					1255
taaa	acca	atc c	ataa	aatt	aļaa	itaac	cttta	a ctt	agcg	ıtat	tttt	t				1300
	)> 11 .> 39															

<400> 112 Arg His Arg Asn Arg Ala Arg Cys Thr Arg Lys Gln Ala Arg Asn Asn Lys Ser Phe Arg His Glu Asn Tyr Phe Tyr Lys Val Leu Gly Ser Ala Thr Ser Gln Ile Glu Ser Leu Lys Lys Arg Glu Asn Ala Leu Phe Asp 40 His Leu Asp Ser Leu Lys Ser Leu Leu Glu Lys Thr His Trp Glu Lys 55 60 Glu Lys Phe Thr Pro Pro Ile Asn Glu Lys Glu Leu Asn Arg Gln Leu 70 75 Lys Glu Val Arg Trp Phe Asn Lys Glu Thr Pro Thr Ser Lys Asn Thr 90 Tyr Lys Lys Ile Gln Lys Leu Ala Val Tyr Lys Ser Pro Leu Ile Lys 1.05 Asp Tyr Leu Tyr Thr Ile Lys Lys Leu Phe Ala Thr Gln Lys Lys Ile 120 Ile Asp Leu Glu Lys Asn Tyr Lys Asp Leu Arg Ala Leu Lys Glu Glu 135 140 Phe Ser Lys Asp Leu Glu Thr Asp Leu Ser His Ser Lys Lys Arg Phe 150 155 Glu Leu Tyr Thr Arg Leu Lys Ser Met Ser Lys Val Phe Ile Ser Lys 165 170 Ser Ile Val Lys Asn Leu Glu Lys Ile Ala Leu Asp Phe Lys Ser Asp 185 180 Arg His Ser Ile Ser Gln Arg Ala Phe Glu Phe Phe Lys Tyr Met Asn 200 Tyr Gln Asn Leu Ser Leu Thr Asp Lys Gly Asn Met Phe Leu Val Ala 215 Lys Phe Phe Lys Asp Ser Ala Leu Leu Val Asn Ile Ala Arg Phe Glu 230 235 Met Lys Lys Ile Asp Asp Ser Val Lys Asn Ser Asn Pro Gln Asp Asn 245 250 Leu Leu Asp Lys Gln Val Trp Leu Asn Leu Leu Glu His Leu Lys Arg 265 Leu Glu Glu Glu Asn Tyr Cys Phe Ala Lys Lys Arg Lys Glu Phe Leu 280 Glu Thr Arg Ala Met Glu Leu Ser Lys Asp Leu Lys Phe Leu Thr Gln 295 300 Ala Asn Glu Asn Asp Leu Pro Ile Tyr Glu Arg Gly Gln Arg Asp Lys 310 315 Ile Ile Lys Arg Cys Glu Lys Ser Leu Asn Phe Leu Gln Lys Glu Leu 325 330 Gln Cys Phe Lys Thr Leu Leu Lys Ser Ala Ser Ile Ala Leu Glu Asn 345 Leu Gln Asn Asn His Gln Ile Thr Ala Val Thr Gln Asp Thr Gln Glu 360 Asn Thr Asn Ala Leu Lys Asn Thr Thr Gln Asp Phe Asn Lys Thr Thr 375 Asn Glu Pro Thr Asn Pro Asn Asn Asn Tyr Gly Met Asp Phe 385 390

<213 <213	0> 11 l> 63 2> Di 3> He	0 A N	obact	cer p	pyloi	ri										
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gtt	Met Lys Lys Ile 1															60 116
_														gtg Val		164
	_		_					_		_			_	gaa Glu 35		212
														ttt Phe		260
														ttg Leu		308
														tct Ser		356
														agc Ser	_	404
														ttc Phe 115	gct Ala	452
														ata Ile		500
			aca Thr			taaa	attag	gct a	actt	tgat	t tt	taag	gtaga	a		548
			aagco				agtga	a aac	caato	gttc	aaat	agco	cta a	acggo	taaac	608

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Met His Val His Phe Val Phe Val Thr Lys Tyr Arg Arg Ser Ala Phe
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Asn Lys Glu Val Ile Asp Phe Leu Gly Ser Val Phe Ala Lys Val Cys
                            40
Lys Asp Phe Glu Ser Glu Leu Val Glu Phe Asp Gly Glu Ser Asp His
                        55
                                             60
Val His Leu Leu Ile Asn Tyr Pro Pro Lys Val Ser Val Ser Lys Leu
                    70
                                        75
Val Asn Ser Leu Lys Gly Val Ser Ser Arg Leu Thr Arg Gln His His
                                     90
Phe Lys Ser Val Glu Ala Ser Leu Trp Gly Lys His Leu Trp Ser Pro
                                105
                                                     110
Ser Tyr Phe Ala Gly Ser Cys Gly Asp Ala Pro Leu Glu Met Ile Lys
                            120
Gln Tyr Ile Gln Asp Gln Glu Thr Pro His
    130
                        135
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caaccaccgt gtttttgaaa ccttaaaaga gccttataac atg caa ttt cat tca
                                             Met Gln Phe His Ser
tct agc gcg ttg att acg cct ttt aaa aaa gat ttg agc gtt gat gag
                                                                   163
Ser Ser Ala Leu Ile Thr Pro Phe Lys Lys Asp Leu Ser Val Asp Glu
                 10
gcc gct tat gaa acc ttg atc aag cgc caa att ttt cag ggc atg gac
                                                                   211
Ala Ala Tyr Glu Thr Leu Ile Lys Arg Gln Ile Phe Gln Gly Met Asp
             25
gca tgc gtg cct gtt ggc acg aca gga gaa tcc gcc acg ctc acc cac
                                                                   259
Ala Cys Val Pro Val Gly Thr Thr Gly Glu Ser Ala Thr Leu Thr His
         40
                             45
aaa gag cac atg cgt tgc att gaa atc gcc ata gaa act tgc aaa aac
                                                                   307
Lys Glu His Met Arg Cys Ile Glu Ile Ala Ile Glu Thr Cys Lys Asn
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							cgc Arg									355
							ctt Leu									403
			_				tgc Cys	-	_					-		451
				_		-	cat His 125					-		_		499
-			-	_			gat Asp	-		-	-					547
	_	-			-		aaa Lys			_	_	_				595
	_			_			ggc Gly		_			-		_	-	643
							aaa Lys									691
							ggg Gly 205									739
			_			_	att Ile			_	_		_			787
						_	cta Leu	-				_			_	835
							gaa Glu								_	883
							att Ile	_			-		_	-		931
							acg Thr									979

280 285 290

caa caa tat gag gta att gca tgaatggttc caatcacatg aaaaataaaa 1030 Gln Gln Tyr Glu Val Ile Ala 300 ccctagtgat cagcggcgcg actagaggga ttggcaaggc gatattgtac 1080 <210> 116 <211> 300 <212> PRT <213> Helicobacter pylori <400> 116 Met Gln Phe His Ser Ser Ser Ala Leu Ile Thr Pro Phe Lys Lys Asp Leu Ser Val Asp Glu Ala Ala Tyr Glu Thr Leu Ile Lys Arg Gln Ile Phe Gln Gly Met Asp Ala Cys Val Pro Val Gly Thr Thr Gly Glu Ser 40 Ala Thr Leu Thr His Lys Glu His Met Arg Cys Ile Glu Ile Ala Ile 55 60 Glu Thr Cys Lys Asn Thr Lys Thr Pro Ser Asn Ser Arg Met Lys Val 70 75 Leu Ala Gly Val Gly Ser Asn Ala Thr Ser Glu Ser Leu Ser Leu Ala 85 90 Lys Phe Ala Gln Lys Ile Gly Ala Asp Ala Ile Leu Cys Val Ser Pro 105 Tyr Tyr Asn Arg Pro Thr Gln Gln Gly Leu Phe Glu His Tyr Lys Thr 115 120 125 Ile Ala Gln Ser Val Glu Ile Pro Val Met Leu Tyr Asp Val Pro Ser 135 Arg Thr Gly Val Ser Ile Glu Val Pro Thr Ala Leu Lys Leu Phe Arg 150 155 Glu Val Pro Asn Ile Lys Ala Ile Lys Glu Ala Ser Gly Ser Leu Lys 170 Arg Val Thr Glu Leu His Tyr Tyr Glu Lys Asp Phe Lys Ile Phe Ser 185 Gly Glu Asp Ser Leu Asn His Ser Ile Met Phe Ser Gly Gly Cys Gly 195 200 205 Val Ile Ser Val Thr Gly Asn Leu Met Pro Asn Leu Ile Ser Gln Met 215 220 Val Asn Cys Ala Leu Lys Gln Lys Tyr Gln Gln Ala Leu Glu Ile Gln 235 230 Asn Lys Leu Phe Cys Leu His Gln Ala Leu Phe Val Glu Thr Asn Pro 245 250 Ile Pro Ile Lys Met Ala Met His Leu Ala Gly Leu Ile Glu Asn Pro 265 Ser Tyr Arg Leu Pro Leu Val Ala Pro Ser Lys Glu Thr Ile Gln Leu 280 Leu Glu Lys Thr Leu Gln Gln Tyr Glu Val Ile Ala 295

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ttaaaaa atg ctt ttg atc gtt ttt ttt aaa ttt tat ttt caa tat tca
        Met Leu Leu Ile Val Phe Phe Lys Phe Tyr Phe Gln Tyr Ser
att aaa aaa aaa tca ttt tat ttt att ttt gtt ata att caa gct att
                                                                   157
Ile Lys Lys Ser Phe Tyr Phe Ile Phe Val Ile Ile Gln Ala Ile
 15
                     20
                                                                   205
ttt att ttc aat cta agg agg tgt cgc atg gac aat caa aag ata acg
Phe Ile Phe Asn Leu Arg Arg Cys Arg Met Asp Asn Gln Lys Ile Thr
cat caa aat atc acg caa aaa caa ggc gag ctt aaa aga gac atg aaa
                                                                   253
His Gln Asn Ile Thr Gln Lys Gln Gly Glu Leu Lys Arg Asp Met Lys
atg cgc cat ctc tta atg att gca ttt gga gga gcg atc ggc aca ggg
                                                                   301
Met Arg His Leu Leu Met Ile Ala Phe Gly Gly Ala Ile Gly Thr Gly
ctt ttt gta ggc act ggg ggt aat att gcg agc gct ggc cct tta ggg
                                                                   349
Leu Phe Val Gly Thr Gly Gly Asn Ile Ala Ser Ala Gly Pro Leu Gly
                         85
                                              90
acc ttg atc gct tat tgt ttt gga ggg ctt gtg gtc tat tgt atc atg
                                                                   397
Thr Leu Ile Ala Tyr Cys Phe Gly Gly Leu Val Val Tyr Cys Ile Met
95
                    100
                                        105
ctc tct tta ggc gaa ttg gct agc gtt tat ccc act aca gga agt ttt
                                                                   445
Leu Ser Leu Gly Glu Leu Ala Ser Val Tyr Pro Thr Thr Gly Ser Phe
                115
                                                         125
ggg gat tat gcg gct aaa ttc ata ggc cct ggc acg ggc tat atg gtt
                                                                   493
Gly Asp Tyr Ala Ala Lys Phe Ile Gly Pro Gly Thr Gly Tyr Met Val
            130
                                                     140
ttt tgg atg tat tgg ctt ggc tgg gtg atc acg gtg gcg tta gaa tac
                                                                   541
Phe Trp Met Tyr Trp Leu Gly Trp Val Ile Thr Val Ala Leu Glu Tyr
        145
                            150
atc gct ata ggc atg ctc atg caa cgc tgg ttt gcg gat att ccc atc
                                                                   589
Ile Ala Ile Gly Met Leu Met Gln Arg Trp Phe Ala Asp Ile Pro Ile
                        165
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cat tat tgg gtt att tta tgc att gcg tta gtt ttt tta ttg aac ttt

His 175	Tyr	Trp	Val	Ile	Leu 180	Cys	Ile	Ala	Leu	Val 185	Phe	Leu	Leu	Asn	Phe 190	
					ttt Phe											685
	-				atc Ile	_										733
					tcg Ser								-			781
					ggg Gly											829
_		_		_	gtt Val 260			_								877
		-			gag Glu				_	_	_		_			925
					ttg Leu		-		-							973
					gtt Val				_		_				_	1021
	_				agc Ser	-		-	_			-				1069
					tat Tyr 340											1117
					acc Thr											1165
					tcc Ser											1213
					ggc Gly											1261

Ser Phe Ser Leu 400			att tat gcc aaa gaa aat Ile Tyr Ala Lys Glu Asn 410	1309
			ttc acg gtg att att gtg Phe Thr Val Ile Ile Val 425 430	1357
Trp Val Ser Val			ttc cgc aag caa tac tta Phe Arg Lys Gln Tyr Leu 445	1405
			tat aaa gcc ccc ttt cta Tyr Lys Ala Pro Phe Leu 460	1453
			tgt gcc atc ggc gtg att Cys Ala Ile Gly Val Ile 475	1501
		_	ggg atg att tta acg att Gly Met Ile Leu Thr Ile 490	1549
	_		tat ttt aca caa aaa gct Tyr Phe Thr Gln Lys Ala 505 510	1597
	aaa aaa gat		tcttttc ttaattttga	1644
Asn Glu Asn Asn		Leu Ile		
Asn Glu Asn Asn	Lys Lys Asp 515		aaaa ccctttttat ctctttggct	1704 1709
Asn Glu Asn Asn agtttagcaa atttt	Lys Lys Asp 515 aagga agtaa		aaaa ccctttttat ctctttggct	
Asn Glu Asn Asn  agtttagcaa atttt ttagc  <210> 118 <211> 519 <212> PRT <213> Helicobact  <400> 118	Lys Lys Asp 515 aagga agtaa er pylori	ccatg atgaaa		
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Leu Gly Glu Leu Ala Ser Val Tyr Pro Thr Thr Gly Ser Phe Gly Asp Tyr Ala Ala Lys Phe Ile Gly Pro Gly Thr Gly Tyr Met Val Phe Trp Met Tyr Trp Leu Gly Trp Val Ile Thr Val Ala Leu Glu Tyr Ile Ala Ile Gly Met Leu Met Gln Arg Trp Phe Ala Asp Ile Pro Ile His Tyr Trp Val Ile Leu Cys Ile Ala Leu Val Phe Leu Leu Asn Phe Phe Ser Val Lys Ile Phe Ala Glu Gly Glu Phe Phe Phe Ser Leu Ile Lys Val Leu Ala Val Ile Ala Phe Ile Gly Ile Gly Ala Ile Gly Ile Ile Tyr Gln Ile Tyr Ser His Gly Phe Gly Ser Ile Phe Asp Asn Phe His Phe Gly Asp Lys Gly Phe Phe Pro Asn Gly Ser Ala Ala Val Phe Ser Ala Met Leu Ala Val Ile Phe Ala Phe Thr Gly Thr Glu Val Ile Gly Val Ala Val Gly Glu Thr Lys Asn Ala Ser Glu Val Met Pro Lys Ala Ile Lys Ala Thr Leu Trp Arg Ile Val Phe Phe Phe Leu Gly Ser Val Phe Val Ile Ser Val Phe Leu Pro Met Asn Asp Ser Ser Ile Thr Gln Ser Pro Phe Val Ser Val Leu Glu Arg Ile Asn Leu Pro Phe Ile Gly Met Gly Ile Pro Tyr Val Ala Asp Ile Met Asn Ala Val Ile Ile Thr Ala Met Phe Ser Thr Ala Asn Ser Gly Leu Tyr Gly Ala Ser Arg Met Ile Tyr Gly Leu Ser Lys Gln Lys Met Phe Phe Lys Val Phe Ser Gln Leu Asn Arg Gln Gly Thr Pro Thr Tyr Ala Met Phe Phe Ser Leu Ser Phe Ser Leu Ile Gly Leu Leu Val Gln Ile Tyr Ala Lys Glu Asn Val Val Glu Ala Leu Ile Asn Val Ile Ser Phe Thr Val Ile Ile Val Trp Val Ser Val Ser Val Ser Gln Tyr Ser Phe Arg Lys Gln Tyr Leu Lys Ala Gly His Ser Leu Glu Asp Leu Pro Tyr Lys Ala Pro Phe Leu Pro Phe Leu Gln Leu Ile Gly Ile Thr Gly Cys Ala Ile Gly Val Ile Gly Ser Ala Met Asp Lys Asp Gln Arg Ile Gly Met Ile Leu Thr Ile Val Phe Ala Val Ile Cys Tyr Ile Gly Tyr Tyr Phe Thr Gln Lys Ala Asn Glu Asn Asn Lys Lys Asp Leu Ile 

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taaggcggtg gtgaatatct ccactgaaaa gaagattaaa aacaatttta taggtggcgg 180
tgtgtttaat gacccctttt tccaacaatt ttttgggggat ttgggtggc atg att cct 238
                                                       Met Ile Pro
                                                        1
aaa gaa aga atg gaa agg gct tta ggc agc ggc gta atc att tct aaa
                                                                   286
Lys Glu Arg Met Glu Arg Ala Leu Gly Ser Gly Val Ile Ile Ser Lys
                                                                   334
gac ggc tat att gta act aat aac cat gtg att gat ggc gcg gat aag
Asp Gly Tyr Ile Val Thr Asn Asn His Val Ile Asp Gly Ala Asp Lys
                     25
att aaa gtt acc att cca ggg agc aat aaa gaa tat tcc gcc act cta
                                                                   382
Ile Lys Val Thr Ile Pro Gly Ser Asn Lys Glu Tyr Ser Ala Thr Leu
gta ggc acc gat tct gaa agc gat tta gcg gtg att cgc acc act aaa
                                                                   430
Val Gly Thr Asp Ser Glu Ser Asp Leu Ala Val Ile Arg Ile Thr Lys
             55
                                  60
gac aat ctg ccc acg atc aaa ttc tct gat tct aat gat att tca gtg
                                                                   478
Asp Asn Leu Pro Thr Ile Lys Phe Ser Asp Ser Asn Asp Ile Ser Val
         70
                             75
ggc gat ttg gtt ttt gcg att ggt aac cct ttt ggc gtg ggc gaa agc
                                                                   526
Gly Asp Leu Val Phe Ala Ile Gly Asn Pro Phe Gly Val Gly Glu Ser
     85
gtt acg caa ggc att gtt tca gcg ctc aat aaa agc ggg att ggg atc
                                                                   574
Val Thr Gln Gly Ile Val Ser Ala Leu Asn Lys Ser Gly Ile Gly Ile
100
                    105
                                         110
                                                             115
aac agc tat gag aat ttc att caa aca gac gct tcc atc aat cct gga
                                                                   622
Asn Ser Tyr Glu Asn Phe Ile Gln Thr Asp Ala Ser Ile Asn Pro Gly
                120
                                     125
aat tcc ggc ggc gct tta att gat agc cgt gga ggg tta gtg ggg att
                                                                   670
Asn Ser Gly Gly Ala Leu Ile Asp Ser Arg Gly Gly Leu Val Gly Ile
            135
                                140
aat acc gct att atc tct aaa act ggg ggc aac cac ggc att ggc ttt
                                                                   718
Asn Thr Ala Ile Ile Ser Lys Thr Gly Gly Asn His Gly Ile Gly Phe
```

155

150

					gtt Val 170									766
					ggt Gly									814
_		_	-		tct Ser							_	_	862
					tct Ser									910
					gtc Val									958
		_			ggc Gly 250		_				-	-		1006
					aaa Lys									1054
_	-				aac Asn			-			_			1102
					aac Asn									1150
				_	cgt Arg	_	_	_	_	-		_		1198
					aat Asn 330									1246
					aaa Lys									1294
					tta Leu									1342
					aat Asn									1390

375 380 385

tgataggggt gggtcgttag tcgcatgtct ttgattagag tgaatgggga agcttttaaa 1450 ctctctttag aaagtttaga agaagaccct tttgaaacta aagaaacgct agaaacgct 1510 atcaaacaaa cgagcgttg 1529

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<212> PRT

<213> Helicobacter pylori

<400> 120

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Lys Ser Val 355	Ala Asp Pl	ne Asn His 360		u Lys Tyr Ly: 365	s Gly Lys
Pro Lys Arg 370	Phe Leu Va	ıl Leu Asp 375	Leu Asn Gl	n Gly Tyr Ar	g Ile Ile
Leu Val Lys 385					
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		_		a gat gaa ato n Asp Glu Ilo 20	e Lys Lys
			Tyr His Pr	g gat ttg aat o Asp Leu Asi 35	
				c gcc gct ta n Ala Ala Ty: 50	
	Glu Glu Ly			t cag ttt ggo p Gln Phe Gly 5	
				c aga agc cgt a Arg Ser Arç	
				t ttt ggg aaa e Phe Gly Lys 100	Gly Gly
			Ser Gln Gl	c ttt tct ggd y Phe Ser Gly 115	
				a acc gcc gct e Thr Ala Ala 130	

gtc tct gtt Val Ser Val 135	Leu Asp 7							8
aat aat gag Asn Asn Glu								6
gaa aag att Glu Lys Ile								4
ggc gat ttg Gly Asp Leu 185		Glu Ile				Met Tyr		2
cgc gag aaa Arg Glu Lys 200				_			_	0
gct ctt ttt Ala Leu Phe 215	Gly Gly I							8
acc cta acc Thr Leu Thr								6
aaa gaa aaa Lys Glu Lys								4
tat ttg cag Tyr Leu Gln 265		Leu Ile	-		-	Leu Ser		2
gag ttg aaa Glu Leu Lys 280		-	_		ggaggaa	tcgtgtgc	ga 92	2
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<400> 122								
Met Ser Lys 1	5			10		15		
Gln Asp Glu	20		25			30		
Pro Asp Leu 35	Asn Lys T		Glu Ala 40	Glu Glu	Lys Phe 45	Lys Glu	Ile	

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Asn Ala Ala Tyr Glu Ile Leu Ser Asp Glu Glu Lys Arg Arg Gln Tyr
Asp Gln Phe Gly Asp Asn Met Phe Gly Gly Gln Asn Phe Ser Asp Phe
                                        75
                    70
Ala Arg Ser Arg Gly Pro Ser Glu Asp Leu Asp Asp Ile Leu Ser Ser
Ile Phe Gly Lys Gly Gly Phe Ser Gln Arg Phe Ser Gln Asn Ser Gln
                                105
Gly Phe Ser Gly Phe Asn Phe Ser Asn Phe Ala Pro Glu Asn Leu Asp
                            120
                                                 125
Ile Thr Ala Ala Leu Asn Val Ser Val Leu Asp Thr Leu Leu Gly Asn
                        135
                                            140
Lys Lys Gln Val Ser Ile Asn Asn Glu Thr Phe Ser Leu Lys Ile Pro
                    150
                                        155
Ile Gly Val Glu Glu Gly Glu Lys Ile Arg Val Arg Asn Lys Gly Lys
                                    170
Thr Gly Arg Thr Thr Arg Gly Asp Leu Leu Glu Ile His Ile Glu
            180
                                185
Glu Asp Glu Met Tyr Arg Arg Glu Lys Asp Asp Ile Thr Gln Ile Phe
                            200
Asp Leu Pro Leu Lys Thr Ala Leu Phe Gly Gly Lys Ile Glu Ile Ala
                        215
                                            220
Thr Trp His Lys Thr Leu Thr Leu Thr Ile Pro Pro Asn Thr Lys Ala
                    230
                                        235
Met Gln Lys Phe Arg Ile Lys Glu Lys Gly Ile Lys Asn Arg Lys Thr
                245
                                    250
Ser His Val Gly Asp Leu Tyr Leu Gln Ala Arg Leu Ile Leu Pro Lys
                                265
Thr Glu Thr Leu Ser Asn Glu Leu Lys Ala Leu Leu Glu Lys Glu Leu
        275
                            280
                                                 285
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                                                           Met Ser
ctg act tcg ctt tta aac cca aaa agc cta gaa gat ttt tta ggc caa
                                                                   166
Leu Thr Ser Leu Leu Asn Pro Lys Ser Leu Glu Asp Phe Leu Gly Gln
                             10
gag cat tta gta ggg aaa gac gcc ccc tta ttt aaa gcc cta caa tcc
                                                                   214
Glu His Leu Val Gly Lys Asp Ala Pro Leu Phe Lys Ala Leu Gln Ser
     20
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							ttc Phe									262
							gcc Ala									310
				_	_		aaa Lys			_	_	-		_		358
							tta Leu 90			_	_			-	~	406
		_	_				caa Gln		_						_	454
_		_		-			tta Leu		-	_	_		_			502
	-		_				cga Arg		-	-				-		550
				_	_	-	tta Leu	_			_	_		_		598
	_						gag Glu 170			_						646
							aga Arg							_	_	694
							atc Ile									742
							gga Gly									790
							aaa Lys									838
							cgc Arg 250									886

gaa ttt atc gcc ag Glu Phe Ile Ala Ar 260		l Ile Phe Ala S		
aac gct aac ccg aa Asn Ala Asn Pro As 275				
gtc aaa caa atc gg Val Lys Gln Ile Gl 29	y Tyr Pro Gl			
gtg att tat ctg gc Val Ile Tyr Leu Al 310		-		
atc aat cag gct tt Ile Asn Gln Ala Le 325		l Gln Lys Gly S		
cct aaa cac ctg ct Pro Lys His Leu Le 340		a Lys Asp Tyr I	_	_
tat aac ggc tat gt Tyr Asn Gly Tyr Va 355				-
gtt tct tct caa gg Val Ser Ser Gln Gl 37	y Ile Gly Ph			
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aaaatcaata tttttgg	att gaatt			1350
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Gly Gln Glu His Le	u Val Gly Ly:	s Asp Ala Pro I 25	Leu Phe Lys Ala 30	Leu
Gln Ser Lys His Ph 35	e Pro His Ala 40	a Phe Phe Tyr (	Gly Pro Pro Gly 45	Val
Gly Lys Thr Ser Le	u Ala Gln Ile 55	=		Pro
Ile Leu Leu Phe As 65	n Ala Thr Asp 70	Phe Lys Leu ( 75	Glu Asp Leu Arg	Leu 80

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Lys Leu Lys Asn Tyr Gln Asn Thr Leu Leu Lys Pro Val Val Phe Ile
                85
                                    90
Asp Glu Thr His Arg Leu Asn Lys Thr Gln Gln Glu Phe Leu Leu Pro
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Ile Met Glu Lys Asp His Ala Leu Ile Leu Gly Ala Ser Thr Gln Asp
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                                                 125
Pro Asn Tyr Ser Leu Ser His Ala Ile Arg Ser Arg Ser Phe Ile Phe
                        135
                                            140
Glu Leu Thr Pro Leu Asn Lys Ser Asp Leu Asp Arg Leu Cys Ala Lys
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                                        155
Ala Leu Thr Leu Leu Lys Lys Gln Ile Glu Pro Gly Ala Lys Thr Tyr
                165
                                    170
Leu Leu Asn Asn Ser Ala Gly Asp Ala Arg Ala Leu Leu Asn Leu Leu
                                185
Asp Leu Ser Ala Lys Ile Glu Asp Pro Ile Thr Leu Lys Thr Leu Gln
                            200
                                                 205
Ser Leu Arg Pro His Ser Leu Asn Asp Gly Ser Tyr Ser Asp Asp Thr
                        215
                                             220
His Tyr Asn Leu Thr Ser Ala Leu Ile Lys Ser Leu Arg Gly Ser Asp
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                                        235
Glu Asn Ala Ser Ile Tyr Tyr Leu Ala Arg Leu Ile Ala Gly Gly Glu
                245
                                    250
Asn Pro Glu Phe Ile Ala Arg Arg Leu Val Ile Phe Ala Ser Glu Asp
                                265
Ile Gly Asn Ala Asn Pro Asn Ala Leu Asn Leu Ala Ala Ser Cys Leu
        275
                            280
Phe Ala Val Lys Gln Ile Gly Tyr Pro Glu Ala Arg Ile Ile Leu Ser
                        295
Gln Cys Val Ile Tyr Leu Ala Cys Ser Pro Lys Ser Asn Thr Ala Tyr
                    310
                                        315
Arg Ala Ile Asn Gln Ala Leu Asp Cys Val Gln Lys Gly Ser Leu Tyr
                325
                                    330
Pro Ile Pro Lys His Leu Leu Pro Asn Ala Lys Asp Tyr Leu Tyr Pro
            340
                                345
His Asp Tyr Asn Gly Tyr Val Lys Gln Asp Tyr Leu Glu Lys Pro Leu
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                                                365
Asp Leu Val Ser Ser Gln Gly Ile Gly Phe Glu Lys Thr Leu Leu Glu
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Trp Leu Asp Lys Ile Arg Asn
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                 Met Gly Phe Tyr Ala Gly Leu Asn Ala Leu Asp Tyr
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aac att ggg tat caa tat gtg ttc taaaaaaata agaaatctca ttttatgctt

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Asn Ile Gly Tyr Gln Tyr Val Phe 225
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                            40
Gly Leu Ile Leu Gly Ile Asn Thr Gly Phe Phe Tyr Asn His Glu Ile
                        55
Met Val Arg Thr Asn Ser Ser Ser Ile Thr Gly Asn Ile Leu Asn Tyr
                    70
                                         75
Leu Phe Ala Tyr Gly Leu Arg Phe Gly Tyr Gln Thr Phe Arg Pro Ser
                85
                                     90
Phe Phe Ala Arg Leu Val Lys Pro Asn Ile Ile Gly Arg Arg Ile Tyr
            100
                                105
                                                     110
Ile Gln Tyr Tyr Gly Gly Ala Pro Lys Lys Ala Gly Phe Gly Asp Val
        115
                            120
Gly Phe Gln Ser Val Met Leu Asn Gly Asp Phe Leu Leu Asp Phe Pro
                        135
Leu Pro Phe Val Gly Lys Tyr Leu Tyr Met Gly Gly Tyr Met Gly Leu
145
                    150
                                         155
Gly Leu Gly Val Val Ala His Gly Val Asn Tyr Thr Ala Glu Trp Gly
                                    170
Met Ser Phe Asn Ala Gly Leu Ala Leu Thr Val Leu Glu Lys Asn Arg
                                185
                                                     190
Ile Glu Phe Gly Phe Lys Ile Leu Asn Asn Phe Pro Phe Leu Gln Ser
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                                                 205
Asn Ser Ser Lys Glu Thr Trp Trp Gly Ala Met Ala Asn Ile Gly Tyr
                        215
                                             220
Gln Tyr Val Phe
225
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		caa gaa gtg agt tt Gln Glu Val Ser Le 20		160
	_	gaa aag att aaa ag Glu Lys Ile Lys Se 35		208
		cgc caa gaa agt tt Arg Gln Glu Ser Le 50		256
		ggt aaa act tca at Gly Lys Thr Ser Il 65		304
		atc aag atc acc gc Ile Lys Ile Thr Al 8		352
		gcc att ttg acc aa Ala Ile Leu Thr As 100		400
		atc cac cgg ctc ag Ile His Arg Leu Se 115		448
Glu Glu Val Leu		gaa gat ttt agg tt Glu Asp Phe Arg Le 130		496
		acc att aaa att ga Thr Ile Lys Ile As 145		544
		aga gcc gga atg ct Arg Ala Gly Met Le 16	u Ser Asn Pro	592
		ttt aga atg caa tt Phe Arg Met Gln Ph 180		640
		aaa gct gcc gtt aa Lys Ala Ala Val Ly 195		688
Asp Ile Lys Gln		gaa atc gct aaa ag Glu Ile Ala Lys Ar 210		736

Thr Pro Arg Ile Ala Leu Arg Leu Leu Lys Arg Val Arg Asp Phe Ala 220 225 230	784
cta gtc aaa aat tca agc ttg atg gat tta aac atc act ttg cat gct Leu Val Lys Asn Ser Ser Leu Met Asp Leu Asn Ile Thr Leu His Ala 235 240 245	832
ttg aat gaa tta ggc gtg aat gaa tta ggc ttt gat gaa gcg gat ttg Leu Asn Glu Leu Gly Val Asn Glu Leu Gly Phe Asp Glu Ala Asp Leu 250 255 260	880
gcg tat tta tct ttg ttg gct aac gct caa gga aag ccg gtg ggt ttg Ala Tyr Leu Ser Leu Leu Ala Asn Ala Gln Gly Lys Pro Val Gly Leu 265 270 275 280	928
aac acg att gca gca tct atg aga gaa gat gaa ggc acg att gaa gac Asn Thr Ile Ala Ala Ser Met Arg Glu Asp Glu Gly Thr Ile Glu Asp 285 290 295	976
gtg att gag cct ttt tta ctc gct aat ggt tat tta gag cgc acc gct Val Ile Glu Pro Phe Leu Leu Ala Asn Gly Tyr Leu Glu Arg Thr Ala 300 305 310	1024
aaa ggc aga atc gcc acg cct aaa acc cat gag ctc tta aaa atc ccc Lys Gly Arg Ile Ala Thr Pro Lys Thr His Glu Leu Leu Lys Ile Pro 315 320 325	1072
act tta aac ccc caa act tta ttt taatcttgtt tagaaagaaa attacactac	1126
Thr Leu Asn Pro Gln Thr Leu Phe 330 335	
	1171
330 335	
330 335  aataacgata aaattttaaa gggtgtaaaa gtagattgtt atgtt  <210> 128 <211> 336 <212> PRT <213> Helicobacter pylori  <400> 128	
aataacgata aaattttaaa gggtgtaaaa gtagattgtt atgtt  <210> 128 <211> 336 <211> 336 <212> PRT <213> Helicobacter pylori  <400> 128  Met Lys Glu Arg Ile Val Asn Leu Glu Thr Leu Asp Phe Glu Ile Ser 1 5 10 15	
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Ile His Arg Leu Ser Pro Ala Ile Glu Glu Val Leu Tyr Pro Ala Met
                                                 125
        115
                            120
Glu Asp Phe Arg Leu Asp Ile Ile Ile Gly Ser Gly Pro Ala Ala Gln
                        135
                                             140
Thr Ile Lys Ile Asp Leu Pro Pro Phe Thr Leu Ile Gly Ala Thr Thr
                   150
                                        155
Arg Ala Gly Met Leu Ser Asn Pro Leu Arg Asp Arg Phe Gly Met Ser
                165
                                    170
Phe Arg Met Gln Phe Tyr Asn Pro Ser Glu Leu Ala Leu Ile Ile Lys
                                185
Lys Ala Ala Val Lys Leu Asn Gln Asp Ile Lys Gln Glu Ser Ala Asp
                            200
Glu Ile Ala Lys Arg Ser Arg Gly Thr Pro Arg Ile Ala Leu Arg Leu
                        215
Leu Lys Arg Val Arg Asp Phe Ala Leu Val Lys Asn Ser Ser Leu Met
                    230
                                        235
Asp Leu Asn Ile Thr Leu His Ala Leu Asn Glu Leu Gly Val Asn Glu
                                    250
                245
Leu Gly Phe Asp Glu Ala Asp Leu Ala Tyr Leu Ser Leu Leu Ala Asn
                                265
Ala Gln Gly Lys Pro Val Gly Leu Asn Thr Ile Ala Ala Ser Met Arg
                            280
                                                285
Glu Asp Glu Gly Thr Ile Glu Asp Val Ile Glu Pro Phe Leu Leu Ala
                        295
Asn Gly Tyr Leu Glu Arg Thr Ala Lys Gly Arg Ile Ala Thr Pro Lys
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                                        315
Thr His Glu Leu Leu Lys Ile Pro Thr Leu Asn Pro Gln Thr Leu Phe
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                                                         Met Phe
gaa gat tta aaa ccg cat tta cag gaa tta aga aag cgt ttg atg gtt
                                                                   164
Glu Asp Leu Lys Pro His Leu Gln Glu Leu Arg Lys Arg Leu Met Val
                             10
tct gta gga acg att cta gtg gcg ttt ttg ggg tgc ttt cat ttt tgg
                                                                   212
Ser Val Gly Thr Ile Leu Val Ala Phe Leu Gly Cys Phe His Phe Trp
     20
                         25
                                             30
aaa agt att ttt gaa ttt gtt aaa aat tcc tat aaa ggc acg ctc att
                                                                   260
Lys Ser Ile Phe Glu Phe Val Lys Asn Ser Tyr Lys Gly Thr Leu Ile
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35					40					45					50	
														agt Ser 65		308
														tta Leu		356
			•					_		_			-	att Ile	_	404
		-												gcg Ala		452
														act Thr		500
														gtg Val 145		548
														ttg Leu		596
														gcg Ala		644
														gta Val		692
gcc Ala 195	att Ile	atc Ile	act Thr	ccc Pro	cct Pro 200	gat Asp	gtg Val	gtg Val	agt Ser	caa Gln 205	atc Ile	ttt Phe	atg Met	gcg Ala	ttg Leu 210	740
														atg Met 225		788
														aac Asn		836
							aag Lys 250				tagt	tgaa	ag a	aattt	gattt	889

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Phe Trp Lys Ser Ile Phe Glu Phe Val Lys Asn Ser Tyr Lys Gly Thr
                            40
Leu Ile Gln Leu Ser Pro Ile Glu Gly Val Met Val Ala Val Lys Ile
                        55
Ser Phe Ser Ala Ala Ile Val Ile Ser Met Pro Ile Ile Phe Trp Gln
                    70
                                        75
Leu Trp Leu Phe Ile Ala Pro Gly Leu Tyr Lys Asn Glu Lys Lys Val
                85
                                    90
Ile Leu Pro Phe Val Phe Phe Gly Ser Gly Met Phe Leu Ile Gly Ala
                                105
Ala Phe Ser Tyr Tyr Val Val Phe Pro Phe Ile Ile Glu Tyr Leu Ala
                            120
                                                125
Thr Phe Gly Ser Asp Val Phe Ala Ala Asn Ile Ser Ala Ser Ser Tyr
                        135
Val Ser Phe Phe Thr Arg Leu Ile Leu Gly Phe Gly Val Ala Phe Glu
                    150
                                        155
Leu Pro Val Leu Ala Tyr Phe Leu Ala Lys Val Gly Leu Ile Thr Asp
                                    170
                165
Ala Ser Leu Lys Ala Tyr Phe Lys Tyr Ala Ile Val Val Ile Phe Ile
                                185
Val Ala Ala Ile Ile Thr Pro Pro Asp Val Val Ser Gln Ile Phe Met
                            200
                                                205
Ala Leu Pro Leu Val Gly Leu Tyr Gly Leu Ser Ile Leu Ile Ala Lys
                        215
                                            220
Met Val Asn Pro Ala Pro Lys Asp Asn Glu Asn Asn Asn Glu Asn Asn
                   230
                                        235
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ttt atg cgc cct tta aac acc aac aag ctt tta caa ggg gct tca atc Phe Met Arg Pro Leu Asn Thr Asn Lys Leu Leu Gln Gly Ala Ser Ile 25 30 35 40	9
ctt caa ggc tat gaa gtg aat cct aaa aac gat tgg gct tat tct agg Leu Gln Gly Tyr Glu Val Asn Pro Lys Asn Asp Trp Ala Tyr Ser Arg 45 50 55	
tat tat ttc ttt ata gat tat ggc aat gtg ctt ttt aat aat gac tct Tyr Tyr Phe Phe Ile Asp Tyr Gly Asn Val Leu Phe Asn Asn Asp Sei 60 65 70	
act tta caa gcg aac atg ttc act tat ggg gtg gga ggg gat ttt atg Thr Leu Gln Ala Asn Met Phe Thr Tyr Gly Val Gly Gly Asp Phe Met 75 80 85	
gtc gcc tac gct aaa aac cct atc aac cgc tgg gct ttt ttc ttt ggc Val Ala Tyr Ala Lys Asn Pro Ile Asn Arg Trp Ala Phe Phe Gly 90 95 100	
ttg caa ctg gcc gct aac aca tgg ata ctc aac aat aaa gtc aaa gat Leu Gln Leu Ala Ala Asn Thr Trp Ile Leu Asn Asn Lys Val Lys As 105 110 115 120	)
ttg gtg gtg aat act tgg gat tca tta aaa gat ttc aat ttt cac aac Leu Val Val Asn Thr Trp Asp Ser Leu Lys Asp Phe Asn Phe His Asr 125 130 135	
act tat ttc agg gct att ggg aag ttt ggg gtg cag ttt cgc acg atc Thr Tyr Phe Arg Ala Ile Gly Lys Phe Gly Val Gln Phe Arg Thr Ile 140 145 150	
gtt ttg tat cat aag gtg gat gta gaa att ggc atg aaa atc ttt cta Val Leu Tyr His Lys Val Asp Val Glu Ile Gly Met Lys Ile Phe Leu 155 160 165	
act cct gaa agg cgc agt ttg ttt gaa agg agc ttt ttg ttt ttt gtt Thr Pro Glu Arg Arg Ser Leu Phe Glu Arg Ser Phe Leu Phe Phe Val 170 175 180	
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Lys Leu Gln Gly Ala Ser Ile Leu Gln Gly Tyr Glu Val Asn Pro
                            40
Lys Asn Asp Trp Ala Tyr Ser Arg Tyr Tyr Phe Phe Ile Asp Tyr Gly
                                            60
Asn Val Leu Phe Asn Asn Asp Ser Thr Leu Gln Ala Asn Met Phe Thr
                                        75
                    70
Tyr Gly Val Gly Gly Asp Phe Met Val Ala Tyr Ala Lys Asn Pro Ile
                85
                                    90
Asn Arg Trp Ala Phe Phe Gly Leu Gln Leu Ala Ala Asn Thr Trp
                                105
Ile Leu Asn Asn Lys Val Lys Asp Leu Val Val Asn Thr Trp Asp Ser
                            120
Leu Lys Asp Phe Asn Phe His Asn Thr Tyr Phe Arg Ala Ile Gly Lys
                        135
                                            140
Phe Gly Val Gln Phe Arg Thr Ile Val Leu Tyr His Lys Val Asp Val
                    150
                                        155
Glu Ile Gly Met Lys Ile Phe Leu Thr Pro Glu Arg Arg Ser Leu Phe
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                                    170
Glu Arg Ser Phe Leu Phe Phe Val Ser His Ser Trp His Phe
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                                                                   102
Phe Val Glu Gly Pro Ser Asp Lys Val Phe Leu Glu Val Tyr Leu Tyr
     10
                         15
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                                                                   150
Phe Leu Glu Arg Phe Pro Ile Lys Asn Phe Lys Val Gln Asn Val Asp
 25
                     30
                                         35
                                                              40
gga aaa gat aac ctg tct aaa cga ttg ctt gaa att gaa aaa tac gat
                                                                   198
Gly Lys Asp Asn Leu Ser Lys Arg Leu Leu Glu Ile Glu Lys Tyr Asp
                 45
                                                          55
                                     50
aaa aca ctt atc att ttt gat gcg gat aaa gac tat gag agt aat aaa
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Lys Thr Leu Ile Ile Phe Asp Ala Asp Lys Asp Tyr Glu Ser Asn Lys
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gaa ( Glu (																342
acc of Thr I 105																390
gaa a Glu s	_		_	_	_			_		-				_		438
aaa a Lys <i>B</i>			_		-				-			-				486
tta q Leu (																534
aaa a Lys I																582
aaa g Lys <i>P</i> 185																630
ctt a																678
gca g Ala (												taat	ttaa	aca		724
aaata ggttt													jct g	gtgaa	attgt	784 830
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1 Val E				5					10		=			15	_	

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Glu Ser Lys Gln Thr Ile Ser Glu Glu Gln Ile Phe Leu Phe Pro Asn
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Asn Gln Asp Asp Gly Asp Leu Glu Thr Leu Leu Leu Lys Ile Ala Asn
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His Lys Glu Phe Ile Asn Cys Phe Glu Ser Tyr Leu Asp Cys Ile Lys
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Lys Lys Glu His Tyr Lys Pro Ile Lys Asn Ile Arg Lys Ser Lys Trp
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Tyr Ala Tyr Leu Glu Ala Leu Gly Leu Glu Lys Phe Phe Gln Tyr Thr
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Trp Asp Thr Lys Lys Lys Asn Asn Lys Lys Leu Ile Ile Asp Asp
               165
                                  170
Lys Asp Gly Asp Glu Ile Glu Ile Lys Asp Gln Tyr Lys Gly Asp Tyr
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Glu Glu Leu Lys Lys Val Leu Asp Leu Asn Ser Lys Ser Leu Ile Pro
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                                                                158
Val Trp Asp Gly Asn Thr Ala Ser Ser Asn Ala Leu Arg Gln Ala Gln
att gat gtc atc gca gcc tat cct atc acc cca tca acg ccc att gtg
                                                                206
Ile Asp Val Ile Ala Ala Tyr Pro Ile Thr Pro Ser Thr Pro Ile Val
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                            35
caa aat tat ggc tcg ttt aag gat aat ggc tat gtt gat ggc gaa ttc
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Gln Asn Tyr Gly Ser Phe Lys Asp Asn Gly Tyr Val Asp Gly Glu Phe
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					gtc Val										350
					tta Leu										398
	-	_		_	aat Asn	_	-								446
					atg Met 130										494
_		-	_		ccc Pro		_	-		-				_	542
					cat His										590
					tgc Cys							_	-		638
					tac Tyr					_				_	686
					gat Asp 210		_	_	_					-	734
					gag Glu										782
					att Ile										830
					cat His						_		_	_	878
					gcg Ala										926

Val Ala Ala 285		atg cgt Met Arg 290	l rys r		Ile L					974
acc atc cat Thr Ile His 300	_	_			-	_		_	_	1022
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ggc act atg Gly Thr Met		_	Asn G	_	_		_			1118
acg caa ggg Thr Gln Gly 350										1166
ggc gaa agg Gly Glu Arg 365	-	_	Ala H		Cys G			_	_	1214
atc aat gaa Ile Asn Glu 380										1262
ttc gta ggc Phe Val Gly				-			ıaagg	jaa		1308
	Leu His 400	Gly Pro	Lys Me	et Ser 405	Phe Pl	he	ıaagg	jaa		1308
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Asn Pro Gln Glu Ala Tyr Asp Phe Thr Leu Met Ala Phe Arg Ile Ala
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                    150
Glu His Gln Lys Val Arg Val Pro Thr Ile Val Asn Gln Asp Gly Phe
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Leu Cys Ser His Thr Val Gln Asn Val Arg Pro Leu Ser Asp Ala Val
                                185
Ala Tyr Gln Phe Val Gly Glu Tyr Gln Thr Lys His Ser Leu Leu Asp
                            200
Phe Asp Lys Pro Val Ser Tyr Gly Ala Gln Ala Glu Glu Glu Trp His
                        215
                                            220
Tyr Glu His Lys Ala Gln Leu His His Ala Ile Met Ser Ala Ser Ser
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                                        235
Val Ile Glu Glu Val Phe Asn Asp Phe Ala Lys Leu Thr Gly Arg Gln
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Tyr His Leu Thr Lys Thr Phe Gln Leu Glu Asp Ala Glu Ile Ala Ile
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Phe Ala Leu Gly Thr Thr Tyr Glu Ser Ala Ile Val Ala Ala Lys Glu
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                                                285
Met Arg Lys Lys Gly Ile Lys Ala Gly Val Ala Thr Ile His Ser Leu
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Arg Pro Phe Pro Tyr Glu Arg Leu Gly Gln Asp Leu Lys Asn Leu Lys
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Ala Leu Ala Ile Leu Asp Lys Ser Ser Pro Ala Gly Thr Met Gly Ala
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Met Phe Asn Glu Val Thr Ser Ala Val Tyr Gln Thr Gln Gly Thr Lys
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His Pro Val Val Ser Asn Tyr Ile Tyr Gly Leu Gly Glu Arg Asp Met
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                                                365
Thr Ile Ala His Leu Cys Glu Ile Phe Glu Glu Ile Asn Glu Asp Ala
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Val Asn Arg Ala Leu Ala Ala Pro Leu Asn Ile His Gly Asp His Ser

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	_			-		tcg Ser 55				_						192
						aac Asn										240
						att Ile										288
		_		_		aat Asn									-	336
			_		_	gag Glu	_									384
						gcc Ala 135										432
						tac Tyr										480
					-	ttt Phe		-				_			-	528
ttt Phe	gct Ala	gga Gly	gtg Val 180	tat Tyr	gac Asp	aaa Lys	gtg Val	agc Ser 185	gat Asp	gcg Ala	tta Leu	ttg Leu	tat Tyr 190	caa Gln	gcc Ala	576
						ggc Gly										624
						gtt Val 215										672
ggc																720

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768
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Phe Leu Glu Tyr Gly Ile Asn Ser Leu Leu Tyr Gln Val Asp Phe Arg
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cgc aat tac tcg gtt tat ttc aac tac act tat att ttt taagcgatag
Arg Asn Tyr Ser Val Tyr Phe Asn Tyr Thr Tyr Ile Phe
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Thr Lys Val Asp Tyr Thr Phe Phe Asn Lys Lys Val Val Ser Pro Ile
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Lys Arg Tyr Ala Asp Arg Ser Ala Phe Tyr Leu Gly Leu Gly Tyr Gln
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Leu Gly Ser Ile Gln His Asn Ser Ser Asn Leu Asn Leu Ser Gln Gln
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Phe Asn Lys Ser Gln Ile Ile Phe Ser Asp Ser Leu Ser Pro Val Phe
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Lys Asn Ser Tyr Val Ser Asn Gly Leu Gly Val Gln Val Gly Tyr Lys
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                                                     110
Trp Val Gly Lys His Glu Glu Thr Lys Trp Phe Gly Phe Arg Trp Gly
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Leu Phe Tyr Asp Leu Ser Ala Ser Leu Tyr Gly Gln Lys Glu Ser Gln
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Ser Val Ile Ile Ser Thr Tyr Gly Thr Tyr Met Asp Leu Leu Leu Asn
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Ala Tyr Asn Gly Asp Lys Phe Phe Ala Gly Phe Asn Leu Gly Ile Ala
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Phe Ala Gly Val Tyr Asp Lys Val Ser Asp Ala Leu Leu Tyr Gln Ala
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                                                     190
Leu Leu Asp Thr Phe Gly Gly Lys Val Asp Pro Asn Gly Phe Gln
        195
                            200
                                                205
Phe Leu Val Asn Leu Gly Val Arg Leu Gly Asn Lys His Asn Gln Phe
                        215
                                            220
Gly Phe Gly Ile Lys Ile Pro Thr Tyr Tyr Phe Asn His Tyr Tyr Ser
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                                        235
Met Asn Asn Ile Ser Asn Asn Ser Glu Asp Val Leu Lys Val Leu Arg
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gcc ctt tct agc Ala Leu Ser Ser 40		Ala Ser Glu Il											
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gcg gtt atc cat Ala Val Ile His 90	Leu Ala Asn G												
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gtt cct gga acg Val Pro Gly Thr			t Leu Lys Lys '										
act aaa aaa cta	tcg caa ctc a	att gat cct gc	c att aaa ttg	gct gaa 535									

Arg Asn Tyr Ser Val Tyr Phe Asn Tyr Thr Tyr Ile Phe 275 280 285

Thr	Lys	Lys	Leu 155	Ser	Gln	Leu	Ile	Asp 160	Pro	Ala	Ile	Lys	Leu 165	Ala	Glu	
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					aaa Lys		_		_							631
				_	tat Tyr 205		_		_	_		_			_	679
	_	_		_	aat Asn				_			_				727
					gct Ala											775
					aaa Lys	_	_		-	_						823
					ggg ggg	-		_			_				_	871
					gga Gly 285		-		_		_				_	919
					tta Leu											967
					gaa Glu											1015
					gct Ala											1063
	Asn				gcc Ala											1111
	345															

		-			acg Thr			-	 _					1207
_	~	_	~		tac Tyr			_			_	_	-	1255
_		_		_	gga Gly		_							1303
					aat Asn									1351
				_	gcc Ala 445	_	_		_		_	_		1399
					aac Asn									1447
					atc Ile									1495
					aat Asn									1543
					cct Pro				-	_			_	1591
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Phe Leu Val Val Gly Ser Pro Gly Gly Ser Arg Ile Ile Thr Thr Val
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Leu Gln Val Ile Ser Asn Val Ile Asp Tyr Asn Met Asn Ile Ser Glu
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Arg Ile Glu Lys Phe Gly Met Pro Ala Asp Val Lys Asp Asn Leu Thr
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Lys Met Gly Tyr Gln Ile Val Thr Lys Pro Val Met Gly Asp Val Asn
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Asp Ser Asn Ile Ser Lys Glu Pro Leu Glu Phe Ile Ile Gly Thr Asn
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Gln Ile Ile Ala Gly Leu Glu Lys Ala Val Leu Lys Ala Gln Ile Gly
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Glu Trp Glu Glu Val Val Ile Ala Pro Glu Glu Ala Tyr Gly Val Tyr
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Glu Ser Ser Tyr Leu Gln Glu Val Pro Arg Asp Gln Phe Glu Gly Ile
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					att Ile												447
					ccg Pro												495
					agg Arg 145												543
					aca Thr												591
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	Ser			Pro	Leu				Ile					Ile	Ile	Ala	
	Gly				Ala										Glu	Glu	
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	Thr	Gly	Cys	Cys	Gly 165	Gly	His	Gly	Gly	His 170	Gly	Gly	Lys	Lys	Gly 175	Gly	ı

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tga									339
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Glu Arg Ser	20			25			30		
Arg Lys Phe	nas ern	wid 2	ст тт6	тте 2	ser AgT	cys GIU	тте ;	ser A	7211

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tac acg ctc a Tyr Thr Leu I 185	Lys Ile Arg (				
tcc att ctc o Ser Ile Leu A 200			-		
ttt tcc atg o		[le Gly Pro			
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cca gtt att a Pro Val Ile 1 250			Asp Leu Phe		
acg ctt gca c Thr Leu Ala C 265	Gly Ile Gly H				
aaa aac aag g Lys Asn Lys 0 280					
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tcc att gcc gat ggc tta gcc ttt ggc ttt ttg tct tat agt atc atc
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Ser Ile Ala Asp Gly Leu Ala Phe Gly Phe Leu Ser Tyr Ser Ile Ile
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                                400
aaa ttg gtt caa aaa gac ttc aaa gca ctc aat tca ggc att atc att
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Lys Leu Val Gln Lys Asp Phe Lys Ala Leu Asn Ser Gly Ile Ile Ile
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                            415
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ctc tgc atc att tct gtt tct gta ttt atc ttt cgt taagctcttt
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Leu Cys Ile Ile Ser Val Ser Val Phe Ile Phe Arg
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Phe Arg Ala Gly Leu Thr Thr Phe Ile Thr Met Ile Tyr Ile Val Pro
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Leu Ser Ala Thr Ala Ile Ile Thr Ile Leu Ser Ser Val Phe Asn Gly
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Leu Trp Ala Asn Thr Pro Ile Ala Met Ser Val Gly Leu Gly Leu Ser
                    70
Ala Tyr Phe Ser Phe Gly Leu Val Gln Gly Leu Lys Leu Pro Trp Gln
                85
                                    90
Ser Ala Leu Gly Ile Val Ala Leu Ser Gly Ala Ile Phe Val Ile Leu
                                105
Ser Phe Thr Lys Phe Arg Ser Trp Val Met Arg Ser Ile Pro Ser Asp
       115
                            120
                                                125
Leu Arg Arg Ala Val Ser Ala Gly Ile Gly Ala Phe Ile Ala Phe Ile
   130
                        135
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Gly Leu Lys Glu Met His Ile Val Val Thr His Xaa Ala Thr Leu Val
145
                    150
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Thr Leu Gly Asp Phe Gly Asp Pro His Val Leu Leu Gly Val Val Gly
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                                    170
Ile Ile Leu Thr Phe Ala Leu Tyr Thr Leu Lys Ile Arg Gly Ser Phe
                                185
Ile Ile Ala Val Leu Ile Thr Ser Ile Leu Ala Trp Val Leu Lys Leu
                            200
                                                205
Ala Pro Tyr Pro Ser Glu Phe Phe Ser Met Pro Ala Ser Ile Gly Pro
                        215
                                            220
Ile Ala Phe Gln Leu Asp Phe Lys Gly Ile Phe Phe Asp Ala Ser Gly
                    230
                                        235
Ala Phe Thr Leu Ala Leu Val Pro Val Ile Ile Thr Phe Phe Val Thr
                245
                                    250
Asp Leu Phe Asp Ser Leu Gly Thr Leu Ala Gly Ile Gly His Lys Thr
            260
                                265
Asp Phe Phe Asn Asp Glu Glu Lys Asn Lys Glu Leu Glu Lys Thr Leu
                            280
                                                 285
Glu Ala Asp Ala Val Ala Ser Leu Gly Ser Ala Val Val Gly Val Ser
                        295
                                            300
Thr Thr Ala Phe Ile Glu Ser Ala Ser Gly Val Glu Glu Gly Gly
                    310
                                        315
Arg Thr Gly Leu Thr Ala Val Phe Thr Gly Leu Phe Phe Val Leu Thr
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                                    330
Leu Phe Cys Leu Pro Leu Leu Lys Ala Ile Pro Ser Asn Ala Ile Tyr
            340
                                345
Pro Val Leu Val Val Val Gly Val Leu Met Phe Ser Val Leu Glu Gly
                            360
                                                 365
Val Asn Phe Lys Asp Met Ala Ile Ser Val Ser Thr Phe Leu Thr Val
                        375
                                             380
Val Met Met Pro Leu Thr Phe Ser Ile Ala Asp Gly Leu Ala Phe Gly
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                                        395
Phe Leu Ser Tyr Ser Ile Ile Lys Leu Val Gln Lys Asp Phe Lys Ala
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                                    410
Leu Asn Ser Gly Ile Ile Leu Cys Ile Ile Ser Val Ser Val Phe
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Ile Phe Arg
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caacc atg acc cct cac atc aac gcc aaa atc ggc gat ttt tat cct caa 110
     Met Thr Pro His Ile Asn Ala Lys Ile Gly Asp Phe Tyr Pro Gln
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	ctt Leu														158
	tta Leu														206
	agc Ser		_		-		 				_				254
_	ggc Gly 65				_	_		_		_					302
	cag Gln			_						_				_	350
	aaa Lys														398
	tct Ser													-	446
	cct Pro														494
	ggt Gly 145	_													542
	tct Ser														590
	gct Ala														638
	aac Asn														686
	aaa Lys														734
	ata Ile 225								tago	cttt	tt t	geed	ccat	a	784

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Ser Gly Lys Tyr Lys Gly Arg Gly Ile Ser Leu Met Gly His Gly Met
Gly Ile Ala Ser Cys Thr Ile Tyr Val Thr Glu Leu Ile Lys Thr Tyr
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Gln Val Lys Glu Leu Leu Arg Ile Gly Thr Cys Gly Ala Ile Ser Pro
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Lys Val Gly Leu Lys Asp Ile Ile Met Ala Thr Gly Ala Ser Thr Asp
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                                105
Ser Lys Thr Asn Arg Val Arg Phe Leu Asn His Asp Leu Ser Ala Thr
        115
                            120
                                                 125
Pro Asp Phe Glu Leu Ser Leu Arg Ala Tyr Gln Thr Ala Lys Arg Leu
                        135
Gly Ile Asp Leu Lys Val Gly Asn Val Phe Ser Ser Asp Phe Phe Tyr
                    150
                                         155
Ser Phe Glu Thr His Ala Phe Asp Leu Met Ala Lys Tyr Asn His Leu
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                                    170
Ala Ile Glu Met Glu Ala Ala Gly Leu Tyr Ala Thr Ala Met Glu Leu
            180
                                185
Asn Ala Lys Ala Leu Cys Leu Cys Ser Val Ser Asp His Leu Ile Thr
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                                                 205
Lys Glu Ala Leu Ser Pro Lys Glu Arg Val Glu Ser Phe Asp Asn Met
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                                             220
Ile Ile Leu Ala Leu Glu Met Met Ser
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	_	_			ttg Leu					_	_	_			_	159
					ttc Phe											207
_					tat Tyr		-			-					_	255
					aag Lys 65			_				-	_		_	303
					gcc Ala		_			_					_	351
					ata Ile											399
					gcg Ala											447
					gcc Ala											495
Thr 140	Ile	Ser	Met	Ala	att Ile 145	Gly	Pro	Gly	Val	Val 150	Ala	Phe	Leu	Gly	Gly 155	543
Ala	Lys	Trp	Leu	Phe 160	tta Leu	Leu	Thr	Ala	Ile 165	Leu	Thr	Leu	Leu	Ser 170	Leu	591
Leu	Met	Leu	Leu 175	Lys	gtc Val	Lys	Asp	Ala 180	Pro	Lys	Ile	Ser	Tyr 185	Gln	Ile	639
Lys	Asn	Ile 190	Lys	Ala	tac Tyr	Gln	Pro 195	Asn	Ser	Lys	Ala	Leu 200	Tyr	Leu	Leu	687
Tyr	Leu 205	Ser	Ser	Phe	ttt Phe	Glu 210	Lys	Ala	Phe	Met	Thr 215	Leu	Ile	Phe	Val	735
					tta Leu											783

220	225	230		235
tta atc ttg gtg t Leu Ile Leu Val T		-		
gga ata gcg agc g Gly Ile Ala Ser V 255			-	
atg ctt tct ggc g Met Leu Ser Gly V 270		Ile Val Ser 7		
tta gcc gac tct a Leu Ala Asp Ser S 285		Lys Tyr Leu 1		_
ggg gtg gcg ttt t Gly Val Ala Phe F 300		_		_
caa tct tta gcg t Gln Ser Leu Ala S 3				
tta ggg caa ttc a Leu Gly Gln Phe T 335				
gtg agc ggg ggg t Val Ser Gly Gly I 350				
ttg atc gtt gta g Leu Ile Val Val A 365		Ile Trp Gly I		
ctc aac aac cct t Leu Asn Asn Pro S 380			Phe Pro Leu Asp	
tac aat gag gaa c Tyr Asn Glu Glu G 4				
tat gtt aat att a Tyr Val Asn Ile S 415	gc gaa gaa atc er Glu Glu Ile	att att gtg a Ile Ile Val I 420	aaa tat aat tcc Lys Tyr Asn Ser 425	gat 1359 Asp
cac att agc gaa g His Ile Ser Glu G 430				
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Phe His Ser Ser Pro Leu Leu Val Gly Leu Ala Val Gly Gly Ala
                            40
Tyr Leu Thr Gln Ile Val Phe Gln Thr Pro Met Gly Ile Leu Ser Asp
                        55
                                            60
Lys Ile Gly Arg Lys Val Val Val Met Val Cys Leu Leu Phe Leu
                                        75
Ala Gly Ser Leu Val Cys Phe Ile Ala Asn Asp Ile Val Trp Leu Val
                85
                                    90
Ile Gly Arg Phe Ile Gln Gly Met Gly Ala Leu Gly Gly Val Ile Ser
            100
                               105
Ala Met Val Ala Asp Glu Val Lys Glu Glu Glu Arg Thr Lys Ala Met
                            120
                                                125
Ala Ile Met Gly Ala Phe Ile Phe Ile Ser Phe Thr Ile Ser Met Ala
                        135
                                            140
Ile Gly Pro Gly Val Val Ala Phe Leu Gly Gly Ala Lys Trp Leu Phe
                   150
                                        155
Leu Leu Thr Ala Ile Leu Thr Leu Leu Ser Leu Leu Met Leu Leu Lys
                165
                                    170
Val Lys Asp Ala Pro Lys Ile Ser Tyr Gln Ile Lys Asn Ile Lys Ala
                                185
Tyr Gln Pro Asn Ser Lys Ala Leu Tyr Leu Leu Tyr Leu Ser Ser Phe
                            200
                                                205
Phe Glu Lys Ala Phe Met Thr Leu Ile Phe Val Leu Ile Pro Leu Ala
                        215
Leu Val Asn Glu Phe His Lys Asp Glu Ser Phe Leu Ile Leu Val Tyr
                    230
                                        235
Val Pro Gly Ala Leu Leu Gly Val Leu Ser Met Gly Ile Ala Ser Val
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                                    250
Met Ala Glu Lys Tyr Asn Lys Pro Lys Gly Val Met Leu Ser Gly Val
            260
                                265
Leu Leu Phe Ile Val Ser Tyr Leu Cys Leu Phe Leu Ala Asp Ser Ser
                            280
Phe Leu Gly Lys Tyr Leu Trp Leu Phe Ile Val Gly Val Ala Phe Phe
                        295
                                            300
Phe Ile Gly Phe Ala Thr Leu Glu Pro Ile Met Gln Ser Leu Ala Ser
                    310
                                        315
Lys Phe Ala Lys Val His Glu Lys Gly Lys Val Leu Gly Gln Phe Thr
                325
                                    330
Thr Phe Gly Tyr Leu Gly Ser Phe Val Gly Gly Val Ser Gly Gly Leu
                                345
Ser Tyr His His Leu Gly Val Ser Asn Thr Ser Leu Ile Val Val Ala
                            360
Leu Gly Leu Ile Trp Gly Leu Ser Leu Phe Leu Leu Asn Asn Pro Ser
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Lys Gln Lys Asn Val Tyr Phe Pro Leu Asp Ala Tyr Asn Glu Glu Gln
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385
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Phe Glu Thr Leu Glu Asp Lys Ile Ile Glu Trp Tyr Val Asn Ile Ser
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Glu Glu Ile Ile Ile Val Lys Tyr Asn Ser Asp His Ile Ser Glu Glu
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Glu Ile Ile His Leu Ala Gln Asn Phe Arg Lys
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                               Met His Ala Glu Phe Phe Thr Phe
gcg ctc atc atg ctt tta att gtg atg gcc cct tat atg tct aga atc
                                                                   160
Ala Leu Ile Met Leu Leu Ile Val Met Ala Pro Tyr Met Ser Arg Ile
                         15
tct cgt ttg cct atc acg gtt gtg gag att tta ttt gga tct gtt ggg
                                                                   208
Ser Arg Leu Pro Ile Thr Val Val Glu Ile Leu Phe Gly Ser Val Gly
 25
                     30
                                          35
gcg tat gtg ggt ttt att gag cct act aag ggc ttt gaa atc atg tcc
                                                                   256
Ala Tyr Val Gly Phe Ile Glu Pro Thr Lys Gly Phe Glu Ile Met Ser
                 45
                                     50
gaa att ggc ttt ttg ttt tta atg ttt tta tgc ggt ttg gaa gtg gag
                                                                   304
Glu Ile Gly Phe Leu Phe Leu Met Phe Leu Cys Gly Leu Glu Val Glu
             60
att tat ctg ttc aaa aaa tta ggg gtt tct ctt tta aaa cgc att ttt
                                                                   352
Ile Tyr Leu Phe Lys Lys Leu Gly Val Ser Leu Leu Lys Arg Ile Phe
         75
                             80
gct tat ctg ttg att tta tac acg ctt tcg ttt att ctt act ttt agc
                                                                   400
Ala Tyr Leu Leu Ile Leu Tyr Thr Leu Ser Phe Ile Leu Thr Phe Ser
     90
                         95
ctt aat tta gag cct att ttt atg gtg att ttc cct att att agt ttg
                                                                   448
Leu Asn Leu Glu Pro Ile Phe Met Val Ile Phe Pro Ile Ile Ser Leu
                    110
ggc atg atc atg act tta gtc aaa gat tat cgt aaa gag att ttg tgg
                                                                   496
Gly Met Ile Met Thr Leu Val Lys Asp Tyr Arg Lys Glu Ile Leu Trp
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				aaa Lys							544
				gtg Val							592
-	_		-	tta Leu			_				640
	-			atc Ile 190	_	_		 	_		688
				atg Met			-			-	736
		_	_	ctc Leu			_		 		784
				gtt Val							832
				cat His				_			880
				ttt Phe 270							928
				tta Leu							976
				ata Ile							1024
				aaa Lys							1072
				tct Ser							1120
				gcg Ala							1168

1267

365 370 aaa ata ctc aac aaa aaa gct tgaatgaaag cttaagcgtc taaatattta

Lys Ile Leu Asn Lys Lys Ala 380

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245

275

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270

285

250

Glu Leu Ile His Lys Leu Asn Asp Val Gly Phe Gly Phe Phe Val Pro 265

Leu Phe Phe Ile His Val Gly Ser Thr Leu Asp Leu Lys Leu Val Phe

Leu Asn Pro His Leu Ile Leu Gln Gly Ile Leu Ile Val Ile Ala Met 295 Leu Ser Leu His Leu Ile Thr Ser Thr Leu Leu Trp Arg Lys Tyr Phe 310 315 Lys Glu Ala Lys His Leu Phe Ser Phe Ala Leu Gly Ala Ser Met Pro 325 330 Leu Thr Phe Leu Val Thr Thr Ala Ala Val Gly Leu Lys Ala Gln Ala 345 Ile Ser Gln Asn Thr Tyr Tyr Ala Leu Leu Met Ala Ala Ile Phe Glu 360 Gly Val Leu Phe Thr Ile Ala Ile Lys Ile Leu Asn Lys Lys Ala 370 375 <210> 153 <211> 1024 <212> DNA <213> Helicobacter pylori <220> <221> CDS <222> (115)...(921) <400> 153 agttggcaaa aacgcagaga cagtaacgca aaggcaaata aagagactca ttttaaacaa 60 gcgaatgcca ttacaaatat aatcagatca gttggtgggt tttttacaaa gatt atg 1 aag aga gtt aga gaa ctt gta aaa aaa cat ccc gag aaa agc agt gtg 165 Lys Arg Val Arg Glu Leu Val Lys Lys His Pro Glu Lys Ser Ser Val 1.0 15 gca tta gta gta tta acc cat gct gca tgc aag aaa gcg aaa gaa ttg 213 Ala Leu Val Val Leu Thr His Ala Ala Cys Lys Lys Ala Lys Glu Leu 25 gac gat aaa gtc cag gat aaa tcc aaa caa gct gaa aaa gaa aat caa 261 Asp Asp Lys Val Gln Asp Lys Ser Lys Gln Ala Glu Lys Glu Asn Gln atc aat tgg tgg aaa tat tca gga tta aca ata gcg aca agt tta tta 309 Ile Asn Trp Trp Lys Tyr Ser Gly Leu Thr Ile Ala Thr Ser Leu Leu 55 tta gcc gct tgt agt gtt ggt gat att gat aaa cag ata gag tta gaa 357 Leu Ala Ala Cys Ser Val Gly Asp Ile Asp Lys Gln Ile Glu Leu Glu 70 75 caa gaa aaa aag gaa gct gaa aac gct agg gat aga gcg aac aag agt 405 Gln Glu Lys Lys Glu Ala Glu Asn Ala Arg Asp Arg Ala Asn Lys Ser 85 ggg ata gaa ctg gaa cag gaa aaa caa aag acc att aaa gaa caa aaa 453 Gly Ile Glu Leu Glu Gln Glu Lys Gln Lys Thr Ile Lys Glu Gln Lys

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	0 0	Gln Asn Cys Gln	gaa aat cat ggc caa 50 Glu Asn His Gly Gln 125	01
			att gct ata gaa gta 54 Ile Ala Ile Glu Val 145	49
	-	•	acc aat caa acc cct 59 Thr Asn Gln Thr Pro 160	97
-			ccc cac tct caa aga 64 Pro His Ser Gln Arg 175	45
			caa aaa gag tta gaa 69 Gln Lys Glu Leu Glu 190	93
		Ala Ile Ala Lys	caa gtg aat ttt tac 74 Gln Val Asn Phe Tyr 205	41
			cct aga gat ttt aag 78 Pro Arg Asp Phe Lys 225	39
			ata cgc tct aaa gct 83 Ile Arg Ser Lys Ala 240	37
			agc cca cct ttc aac 88 Ser Pro Pro Phe Asn 255	35
		tca aaa aat att : Ser Lys Asn Ile ( 265		31
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Val Ala Leu Val Val Leu Thr His Ala Ala Cys Lys Ala Lys Glu

5

. 25

20

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Glu Gln Glu Lys Lys Glu Ala Glu Asn Ala Arg Asp Arg Ala Asn Lys
                                    90
Ser Gly Ile Glu Leu Glu Gln Glu Lys Gln Lys Thr Ile Lys Glu Gln
                                105
Lys Asp Leu Val Lys Lys Ala Glu Gln Asn Cys Gln Glu Asn His Gly
                            120
Gln Phe Phe Met Lys Lys Leu Gly Ile Lys Gly Gly Ile Ala Ile Glu
                        135
                                             140
Val Glu Ala Glu Cys Lys Thr Pro Lys Pro Ala Lys Thr Asn Gln Thr
                    150
                                        155
Pro Ile Gln Pro Lys His Leu Pro Asn Ser Lys Gln Pro His Ser Gln
                165
                                    170
Arg Gly Ser Lys Ala Gln Glu Leu Ile Ala Tyr Leu Gln Lys Glu Leu
                                185
Glu Ser Leu Pro Tyr Ser Gln Lys Ala Ile Ala Lys Gln Val Asn Phe
                            200
Tyr Arg Pro Ser Ser Val Ala Tyr Leu Glu Leu Asp Pro Arg Asp Phe
                        215
                                            220
Lys Val Thr Glu Glu Trp Gln Lys Glu Asn Leu Lys Ile Arg Ser Lys
                    230
                                        235
Ala Gln Ala Lys Met Leu Gly Asn Glu Lys Pro Thr Ser Pro Phe
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                                    250
Asn Leu Ser Lys Pro Phe Val Arg Ser Lys Asn Ile Cys
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aat ggc gtt ttg gca ggc tat ccg gtg gtg gat ttt aaa gtt acc ctt
                                                                   104
Asn Gly Val Leu Ala Gly Tyr Pro Val Val Asp Phe Lys Val Thr Leu
                             10
tat gat ggg agc tac cat gat gtg gat tct tca gaa atg gcg ttt aaa
                                                                   152
Tyr Asp Gly Ser Tyr His Asp Val Asp Ser Ser Glu Met Ala Phe Lys
atc gct ggc tct atg gcg ttt aaa gaa gcg agt cgc gcg gct aac ccg
                                                                   200
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Ile Ala Gly 35	Ser Met	Ala 40	Phe	Lys	Glu	Ala	Ser 45	Arg	Ala	Ala	Asn	Pro 50	
gtt tta cta Val Leu Lev		Met											248
tac atg ggo Tyr Met Gly													296
aat tot ato Asn Ser Med 8!	Asp Asp	_											344
ccg ttg gtg Pro Leu Val	-												392
caa ggg cg Gln Gly Arg 115													440
cct agc aat Pro Ser Asi		Lys											482
tgatttaatt	ataacgct	ct ct	tatt	ttta	a ggg	gggtg	gtta	tagg	gtgct	igt t	ta		535
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                                                                   104
Leu Gly Asn Pro His Ser Ser Ile Leu Arg Leu Gly Pro Thr Thr Ile
acg ctt ctg cct gaa aaa tcc acg cgc tta cct aaa agg ttt tgc ctg
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Thr Leu Leu Pro Glu Lys Ser Thr Arg Leu Pro Lys Arg Phe Cys Leu
                         25
aaa cgc ccc tgc ttg cct tta atg att tca ctg agc gat ttt aaa ggg
                                                                   200
Lys Arg Pro Cys Leu Pro Leu Met Ile Ser Leu Ser Asp Phe Lys Gly
cgt ttg tta gcc cct tta acc gca tta gtg ctg cgg ccg tta tca aaa
                                                                   248
Arg Leu Leu Ala Pro Leu Thr Ala Leu Val Leu Arg Pro Leu Ser Lys
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age aca tee acg get tet tge aac ate ett ttt tea ttg ege aca atg
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Ser Thr Ser Thr Ala Ser Cys Asn Ile Leu Phe Ser Leu Arg Thr Met
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                                  75
att tot ggc gct cca agc toc att aag cgt ttc aag cgt tgg tta cga
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Ile Ser Gly Ala Pro Ser Ser Ile Lys Arg Phe Lys Arg Trp Leu Arg
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                             90
ttg atg aca cga cga tac aat tca ttc aca tcg ctg act gca aac ttc
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Leu Met Thr Arg Arg Tyr Asn Ser Phe Thr Ser Leu Thr Ala Asn Phe
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                        105
ccg cca tct agc gcg act aaa ggc ctt aaa tcc ggt ggc aat acc ggt
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Pro Pro Ser Ser Ala Thr Lys Gly Leu Lys Ser Gly Gly Asn Thr Gly
115
                    120
aaa acc gtg agc atc atc cat tca ggc cta tta cca gaa ttt aaa aag
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Lys Thr Val Ser Ile Ile His Ser Gly Leu Leu Pro Glu Phe Lys Lys
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ctt tct acc act ttc aaa cgc tta atg agt ttt ttc ttt ttc gca tca
                                                                   536
Leu Ser Thr Thr Phe Lys Arg Leu Met Ser Phe Phe Phe Phe Ala Ser
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gaa ttg gtg tct ttc act tct tct ttc aaa ctc tgc aat aag gtg atc
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Glu Leu Val Ser Phe Thr Ser Ser Phe Lys Leu Cys Asn Lys Val Ile
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Lys Ser Ile Ser Ser Asn Lys Ser Leu Ile Ala Ser Pro Pro Ile Cys
    180
                                                                   681
gct aca aag ccc ctg tct tcg tat ctt cgt gag ata ttt tgatactgct
Ala Thr Lys Pro Leu Ser Ser Tyr Leu Arg Glu Ile Phe
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Cys Leu Lys Arg Pro Cys Leu Pro Leu Met Ile Ser Leu Ser Asp Phe
                            40
Lys Gly Arg Leu Leu Ala Pro Leu Thr Ala Leu Val Leu Arg Pro Leu
                        55
Ser Lys Ser Thr Ser Thr Ala Ser Cys Asn Ile Leu Phe Ser Leu Arg
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                    70
Thr Met Ile Ser Gly Ala Pro Ser Ser Ile Lys Arg Phe Lys Arg Trp
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Leu Arg Leu Met Thr Arg Arg Tyr Asn Ser Phe Thr Ser Leu Thr Ala
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Asn Phe Pro Pro Ser Ser Ala Thr Lys Gly Leu Lys Ser Gly Gly Asn
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Thr Gly Lys Thr Val Ser Ile Ile His Ser Gly Leu Leu Pro Glu Phe
                        135
                                            140
Lys Lys Leu Ser Thr Thr Phe Lys Arg Leu Met Ser Phe Phe Phe
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                                        155
Ala Ser Glu Leu Val Ser Phe Thr Ser Ser Phe Lys Leu Cys Asn Lys
                165
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Val Ile Lys Ser Ile Ser Ser Asn Lys Ser Leu Ile Ala Ser Pro Pro
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Ile Cys Ala Thr Lys Pro Leu Ser Ser Tyr Leu Arg Glu Ile Phe
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	-				cgc Arg 70		_		_					_		240
					ttt Phe											288
	_			_	gaa Glu			_	_		_		_			336
					aat Asn											384
-	-	-	_	_	ttg Leu		_		tgat	tgta	agg (	ctttg	gctto	ca		431
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	Ser	Asn	Tyr 20		Arg	Asn	Tyr	Gln 25		Lys	His	Asn	Leu 30		Tyr	
Trp	Gly	Ala 35	Lys	Asp	Tyr	Leu	Gly 40	Cys	Gly	Ala	Gly	Ala 45		Gly	Cys	
Val	Ala 50	Asn	Glu	Arg	Phe	Phe 55	Ala	Lys	Lys	Leu	Ile 60	Glu	Asn	Tyr	Ile	
Tage	Aen	Pro	T.e.1	Gln	Δτα	Gln	t/a1	Glu	ጥኮኍ	Leu		Lare	Cln	Δαη	Tare	

65 70 75 Arg Leu Glu Lys Leu Phe Leu Gly Leu Arg Cys Val Leu Gly Val Glu 85 90 Leu Ser Phe Leu Asp Glu Asn Lys Val Lys Phe Leu Ile Glu Glu Asn 105 Lys Ala Phe Ile Lys Asn Asn Arg Leu Ile Ala Ser Asp Phe Phe Met 120 125 Ala Asp Glu Met Ala Leu Trp Leu Leu 130 135 <210> 161 <211> 1260 <212> DNA <213> Helicobacter pylori <220> <221> CDS <222> (55)...(1179) <400> 161 gttggtgttt gaaaggctct atgaagaaga attaaggcgc aagggttttt tata atg gtc tct ctc tat tta gaa aac ggg ctt ttt ttg caa gcg caa agt ttt 105 Val Ser Leu Tyr Leu Glu Asn Gly Leu Phe Leu Gln Ala Gln Ser Phe 5 15 ggg gct agc ggc acg caa gcg ggc gag ctt gtt ttt aac act tct atg 153 Gly Ala Ser Gly Thr Gln Ala Gly Glu Leu Val Phe Asn Thr Ser Met 20 25 agc ggt tat caa gaa gtc att agc gac cct agc tat aag ggg caa ttt 201 Ser Gly Tyr Gln Glu Val Ile Ser Asp Pro Ser Tyr Lys Gly Gln Phe 35 40 gtg gtt ttt agc atg cct gag att ggg gtt gtg ggt gct aat tct aaa 249 Val Val Phe Ser Met Pro Glu Ile Gly Val Val Gly Ala Asn Ser Lys 55 gat gat gaa too ttt ttt toa tgo goa ggg gtt tta gog ogo cat tac 297 Asp Asp Glu Ser Phe Phe Ser Cys Ala Gly Val Leu Ala Arg His Tyr 70 aac gaa ttt ttt tct aac tca agg gcg gat ttt agc ttg agc gct tat 345 Asn Glu Phe Phe Ser Asn Ser Arg Ala Asp Phe Ser Leu Ser Ala Tyr 85 90 ttg aaa gag cgt ggc gtt tta ggg gtt tgt ggc gtt gat act agg agt 393 Leu Lys Glu Arg Gly Val Leu Gly Val Cys Gly Val Asp Thr Arg Ser 100 105 ttg att aaa acc tta cgc cat cat ggg tgc tta atg atg gtc gct tcc 441 Leu Ile Lys Thr Leu Arg His His Gly Cys Leu Met Met Val Ala Ser

	115					120					125	
acg	ata	gag	cat	gac	aaa	aac	aag	ctt	gaa	gaa	att	

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					tcc Ser					_	~		_			537
					cgt Arg											585
					aaa Lys											633
					aat Asn											681
					ccg Pro 215											729
					gaa Glu										_	777
_					ttg Leu											825
					ccc Pro											873
					ggc Gly											921
					ccc Pro 295											969
					cac His											1017
					cac His											1065
gtg	cgt	tat	aaa	aac	gct	ccc	att	atc	tct	gtc	cag	cac	cac	cca	gaa	1113

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Ser Ser Pro Gly Pro Lys Glu Ser His Tyr Ile Phe Lys Glu Phe Val
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                                                 45
Phe Val Val Phe Ser Met Pro Glu Ile Gly Val Val Gly Ala Asn Ser
Lys Asp Asp Glu Ser Phe Phe Ser Cys Ala Gly Val Leu Ala Arg His
Tyr Asn Glu Phe Phe Ser Asn Ser Arg Ala Asp Phe Ser Leu Ser Ala
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Tyr Leu Lys Glu Arg Gly Val Leu Gly Val Cys Gly Val Asp Thr Arg
                                105
Ser Leu Ile Lys Thr Leu Arg His His Gly Cys Leu Met Met Val Ala
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                                                125
Ser Thr Ile Glu His Asp Lys Asn Lys Leu Glu Glu Ile Leu Lys Asn
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Ala Pro Lys Ile Ser His Ser Pro Leu Val Ser Ser Val Ser Thr Pro
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                                        155
Lys Ile Thr Thr His Gln Arg Ala Thr Phe Asp Phe Lys Thr Leu Asp
                165
                                    170
Tyr Lys Pro Phe Asp Glu Lys Thr Ser His Lys Ile Ile Ala Val Leu
                                185
Asp Phe Gly Ala Lys Gly Asn Ile Leu Asn Glu Leu Gln Asn Val Gly
                            200
Leu Lys Ala Leu Ile Tyr Pro His His Thr Lys Ala Ser Glu Leu Ile
                        215
                                            220
Lys Ala Tyr Glu Lys Lys Glu Ile Ser Gly Ile Phe Leu Ser Asn Gly
                    230
                                        235
Pro Gly Asp Pro Leu Ser Leu Gln Glu Ile Gly Glu Ile Lys Gln
                245
                                    250
Leu Ile Asn Ala Lys Ile Pro Met Leu Gly Ile Cys Leu Gly His Gln
                                265
                                                    270
Leu Leu Ser Ile Ala Gln Gly Tyr Pro Thr Tyr Lys Leu Lys Phe Gly
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280

His His Gly Ser Asn His Pro Val Lys Asn Leu Lys Thr Asn Ala Val 295 300 Glu Ile Thr Ala Gln Asn His Asn Tyr Cys Val Pro Glu Asp Ile Glu 310 315 Glu Ile Ala Ile Ile Thr His Arg Asn Leu Phe Asp Asn Thr Ile Glu 325 330 Gly Val Arg Tyr Lys Asn Ala Pro Ile Ile Ser Val Gln His His Pro 345 Glu Ser Ser Pro Gly Pro Lys Glu Ser His Tyr Ile Phe Lys Glu Phe 360 Val Glu Leu Leu Lys Asp Phe 370 375 <210> 163 <211> 2790 <212> DNA <213> Helicobacter pylori <220> <221> CDS <222> (116)...(2656) <400> 163 ttctccttta attttggatg tttaaagtat aataaactat ctttttaaaa aaataactta 60 aaagagctat aaaatagcct taaaatacgc gataaaacaa caaaaaggaa taccc atg 1 gat att cgc aac gaa ttt tta caa ttt ttt caa aat aaa ggg cat gcc Asp Ile Arg Asn Glu Phe Leu Gln Phe Phe Gln Asn Lys Gly His Ala gtt tat cct agc atg cct tta gtg cct aat gac gct acc ttg ctt ttt 214 Val Tyr Pro Ser Met Pro Leu Val Pro Asn Asp Ala Thr Leu Leu Phe 25 acc aat gcc ggc atg gtg caa ttt aaa gat att ttt acc ggg att gtg 262 Thr Asn Ala Gly Met Val Gln Phe Lys Asp Ile Phe Thr Gly Ile Val 40 cca cgc cct agc att cct aga gcg gca agc tcg caa ttg tgc atg cgc 310 Pro Arg Pro Ser Ile Pro Arg Ala Ala Ser Ser Gln Leu Cys Met Arg 50 55 gca ggc ggc aag cat aac gat ttg gaa aat gtc ggt tat acc gca agg 358 Ala Gly Gly Lys His Asn Asp Leu Glu Asn Val Gly Tyr Thr Ala Arg 70 75 80 cac cac acg ctt ttt gaa atg cta ggg aat ttc tct ttt ggg gat tat 406 His His Thr Leu Phe Glu Met Leu Gly Asn Phe Ser Phe Gly Asp Tyr 85 90 95 ttc aaa gaa gag atc ttg ttt gcg tgg gaa ttt gta acc aag aat 454 Phe Lys Glu Glu Ala Ile Leu Phe Ala Trp Glu Phe Val Thr Lys Asn

		100					105					110				
											gtg Val 125					502
											cct Pro					550 5
											atg Met					598
											cag Gln					646
											ggc Gly					694
											cgc Arg 205					742
											aca Thr					790
											aat Asn					838
											gag Glu					886
											gta Val					934
gca Ala	aga Arg 275	gcg Ala	gta Val	gca Ala	ttc Phe	ttg Leu 280	ctc Leu	gct Ala	caa Gln	ggg Gly	gtg Val 285	cat His	ttc Phe	aat Asn	aag Lys	982
Glu 290	Gly	Arg	Gly	Tyr	Val 295	Leu	Arg	Arg	Ile	Leu 300	agg Arg	Arg	Ala	Leu	Arg 305	1030
cat His	Gly	tat Tyr	tta Leu	atg Met 310	ggc Gly	ttg Leu	aaa Lys	gaa Glu	gcg Ala 315	ttt Phe	tta Leu	tac Tyr	aaa Lys	gtc Val 320	gtg Val	1078

ggc gtg gtg tgc gag caa ttt gct aac acg cat gcg tat ttg aaa gag 1126

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			_	_	tcg Ser		_	_	_			-			_	1222
	_		_		aaa Lys 375			_		_		-		_		1270
					ttc Phe											1318
				_	gcg Ala	-	_				_	_	_	_		1366
					tct Ser											1414
					att Ile											1462
					gaa Glu 455											1510
					ata Ile											1558
					acc Thr											1606
					ctt Leu											1654
tta Leu	gat Asp 515	aca Thr	aaa Lys	aac Asn	ttt Phe	ttt Phe 520	ggg Gly	ctt Leu	aat Asn	ttt Phe	tca Ser 525	ctc Leu	ctt Leu	gaa Glu	atc Ile	1702
					aaa Lys 535											1750

											act Thr				1798
_	_			-							agt Ser				1846
		_		_	_	_	_		_		tcg Ser		-		 1894
		_	_			-		_	_	_	tta Leu 605	_		_	1942
											atg Met				1990
											gaa Glu				2038
											att Ile	_	_	_	 2086
											ggg Gly				2134
											att Ile 685				2182
											aat Asn				2230
											atc Ile				2278
											gcc Ala				2326
											ttg Leu				2374
											cga Arg 765				2422

770											gaa Glu					2470
											aaa Lys					2518
											ggg Gly					2566
											aat Asn					2614
Leu											tta Leu 845					2656
	ratta	aag t	ttga						-						agagca cacaga	
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- Z I J	- 116		macı													
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Met 1 Ala Phe Val	Asp Val Thr	Ile Tyr Asn 35	Arg Pro 20 Ala	Asn 5 Ser Gly	Glu Met Met	Phe Pro Val	Leu Gln 40	Val 25 Phe	10 Pro Lys	Asn Asp	Asp Ile	Ala Phe 45	Thr 30 Thr	15 Leu Gly	Leu Ile	
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Gly Val Leu Ser Pro Leu Pro Lys Pro Ser Ile Asp Thr Gly Met Gly
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                                            220
Leu Glu Arg Val Gln Ala Leu Leu Glu His Lys Leu Asn Asn Phe Asp
                    230
                                        235
Ser Ser Leu Phe Ala Pro Leu Met Glu Glu Ile Ser Glu Leu Thr Ser
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                                    250
Leu Asp Tyr Ala Ser Glu Phe Gln Pro Ser Phe Arg Val Val Ala Asp
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                                                    270
His Ala Arg Ala Val Ala Phe Leu Leu Ala Gln Gly Val His Phe Asn
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                            280
                                                285
Lys Glu Gly Arg Gly Tyr Val Leu Arg Arg Ile Leu Arg Arg Ala Leu
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                                            300
Arg His Gly Tyr Leu Met Gly Leu Lys Glu Ala Phe Leu Tyr Lys Val
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                                        315
Val Gly Val Val Cys Glu Gln Phe Ala Asn Thr His Ala Tyr Leu Lys
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Glu Ser Lys Glu Met Val Val Lys Glu Cys Phe Glu Glu Glu Glu His
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Phe Leu Glu Thr Leu Glu Ser Gly Met Glu Leu Phe Asn Leu Ser Leu
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Lys His Leu Asn Glu Asn Lys Ile Phe Asp Gly Lys Ile Ala Phe Lys
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Leu Tyr Asp Thr Phe Gly Phe Pro Leu Asp Leu Thr Asn Asp Met Leu
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                                        395
Arg Ser His Gly Ala Cys Ala Asp Met Gln Gly Phe Glu Leu Cys Met
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Gln Glu Gln Val Lys Arg Ser Lys Ala Ser Trp Lys Gly Lys Gln Asn
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Asn Ala Asp Phe Ser Ala Ile Leu Asn Ala Tyr Ala Pro Asn Val Phe
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Val Gly Tyr Glu Thr Thr Glu Cys Ser Ala Lys Val Leu Gly Phe Phe
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Asp Ser Asp Phe Lys Glu Ile Thr Asp Ala Asn Pro Asn Gln Glu Val
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Trp Val Leu Leu Glu Lys Thr Pro Phe Tyr Ala Glu Gly Gly Gly Ala
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                                    490
Ile Gly Asp Arg Gly Ala Leu Phe Lys Asp Asn Gly Glu Val Ala Ile
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                                505
Val Leu Asp Thr Lys Asn Phe Phe Gly Leu Asn Phe Ser Leu Leu Glu
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Ile Lys Lys Ala Leu Lys Lys Gly Asp Gln Val Ile Ala Gln Val Ser
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Asp Glu Arg Phe Glu Ile Ala Lys His His Ser Ala Thr His Leu Leu
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Gln Ser Ala Leu Arg Glu Val Leu Gly Ser His Val Ser Gln Ala Gly
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                                    570
Ser Leu Val Glu Ser Lys Arg Leu Arg Phe Asp Phe Ser His Ala Lys
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                                                    590
Ala Leu Asn Asp Glu Glu Leu Glu Lys Val Glu Asp Leu Val Asn Ala
                            600
Gln Ile Phe Lys His Leu Asn Ser Gln Val Glu His Met Pro Leu Asn
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                                            620
Gln Ala Lys Asp Lys Gly Ala Leu Ala Leu Phe Ser Glu Lys Tyr Ala
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Gly Gly Ile His Val Glu Asn Thr Gly Leu Ile Gly Gly Phe Arg Ile
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Val Lys Glu Ser Gly Val Ser Ser Gly Val Arg Arg Ile Glu Ala Val
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Cys Gly Lys Ala Phe Tyr Gln Leu Ala Lys Glu Glu Asn Lys Glu Leu
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Lys Asn Ala Lys Thr Leu Leu Lys Asn Asn Asp Val Ile Ala Gly Ile
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Asn Lys Leu Lys Glu Ser Val Lys Asn Ser Gln Lys Ala Pro Val Ser
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Met Asp Leu Pro Val Glu Lys Ile His Gly Val Asn Leu Val Val Gly
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Val Val Glu Gln Gly Asp Ile Lys Glu Met Ile Asp Arg Leu Lys Ser
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Lys His Glu Arg Leu Leu Ala Met Val Phe Lys Lys Glu Asn Glu Arg
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                                            780
Ile Thr Leu Ala Cys Gly Val Lys Asn Ala Pro Ile Lys Ala Asn Val
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Trp Ala Asn Glu Val Ala Gln Ile Leu Gly Gly Lys Gly Gly Arg
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Ile Met Val Gly Arg Leu Thr Arg Asn Val Glu Leu Lys Tyr Leu Pro
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Ser Gly Ser Ala Ala Ala Thr Ile Gly Leu Ala Thr Ser Arg Phe
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aaa aaa caa gac ggc acg cta ggc gaa gag gtg tgc ttt ata gat gcg
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Lys Lys Gln Asp Gly Thr Leu Gly Glu Glu Val Cys Phe Ile Asp Ala
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625

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ttg caa ttt a Leu Gln Phe M 1		_	_		-	-	450
caa gat agt a Gln Asp Ser I 120			Ser Asn				498
cat aac gct c His Asn Ala P 135							546
gct tac gct a Ala Tyr Ala L 150		n Leu Gln					594
agc gtg cct g Ser Val Pro G		_		Glu Ile			636
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Thr Ser Arg A:	rg Phe Ly	s Lys Gln 40	Asp Gly	Thr Leu	Gly Glu 45	Glu Val	
Cys Phe Ile A	sp Ala Ar	J Leu Phe 55	Gly Arg	Thr Ala		Ala Asn	
Gln Tyr Leu So	er Lys Gl 70	Ser Ser	Val Leu		Gly Arg	Leu Thr 80	
Tyr Glu Ser T	rp Met As _l 85	Gln Thr	Gly Lys 90		Ser Arg		
Ile Thr Ala A		Gln Phe		Lys Lys	Ser Asp		

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Ala Tyr Ala Gl 145	n Asn Ala 150		Lys Glu	Asn Leu 155	Gln Ala	Gln Pro 160	
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ccc ata tta ct Pro Ile Leu Le		_		_		_	258
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ttt caa gtt tg Phe Gln Val Cy 75							354
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gac gct tta gg Asp Ala Leu Gl 105							450

	ttg Leu															498
	aaa Lys							_		_		_				546
	ttg Leu									-			_			594
	aaa Lys 170															642
	aac Asn															690
gag Glu	ggc Gly	aag Lys	ctc Leu	gct Ala 205	tat Tyr	gat Asp	ttg Leu	gcg Ala	tat Tyr 210	GJA aaa	ttt Phe	tta Leu	acg Thr	ccc Pro 215	ttt Phe	738
	tct Ser															786
	ctc Leu															834
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1	Ser			5					10					15		
Leu	Arg	Phe 35		Gly	His	Tyr	His 40		Ile	Leu	Leu	Pro 45		Glu	Ser	
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Ile Asn Ala Thr Ser Ala Ser Leu His Asn Glu Leu Pro Leu Asn Lys
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Ala Tyr Gly Phe Leu Thr Pro Phe Leu Ser Leu Ala Lys Glu Leu Lys
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Thr Pro Phe Gln Asp Gly Lys Asp Met Leu Ile Tyr Gln Ala Ala Leu
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Tyr Ile Ile Glu Thr Leu Ile Lys Ile Leu Ile Leu Val Ala Val Phe
                         10
tcg gct tta gga ggc ttt gcc act tat att gaa agg aaa gtg tta gcc
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Ser Ala Leu Gly Gly Phe Ala Thr Tyr Ile Glu Arg Lys Val Leu Ala
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Tyr Phe Gln Arg Arg Leu Gly Pro Cys Tyr Val Gly Pro Phe Gly Leu
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                                     45
                                                          50
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Leu Gln Val Ala Ala Asp Gly Ile Lys Leu Phe Thr Lys Glu Asp Ile
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							tcc Ser									343
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							gsc Ala									497
							caa Gln 155									535
							tta Leu									583
							agc Ser									631
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-	_		_				gac Asp	_				-	_			727
							tac Tyr 235									775
			_				tta Leu		-							823
							aac Asn									871
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Pro Ile Ile Ala Met Val Ser Ala Phe Val Ser Met Ala Pro Ile Pro
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                                    90
Phe Phe Pro Asn Phe Thr Leu Phe Gly Tyr Glu Ile Lys Pro Leu Ile
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Ser Asp Ile Asn Ile Gly Phe Leu Phe Phe Leu Ala Val Gly Ser Ala
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Val Val Ser Thr Leu Thr Ile Leu Ala Pro Leu Met Val Val Gly Ser
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Leu Ser Leu Val Glu Ile Asn His Tyr Gln Ser Gly Gly Phe Leu Asp
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Trp Leu Val Phe Lys Gln Pro Leu Ala Phe Val Leu Phe Leu Ile Ala
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Ser Tyr Ala Glu Leu Asn Arg Thr Pro Phe Asp Leu Leu Glu His Glu
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Ala Glu Ile Val Ala Gly Tyr Cys Thr Glu Tyr Ser Gly Leu Lys Trp
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Gly Met Phe Phe Leu Ala Glu Tyr Ala His Leu Phe Ala Phe, Ser Phe
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				_	_	gtg Val	_		-							536
						gaa Glu 170										584
_	-	-			_	cta Leu			-	-						632
		_	_			ttt Phe			_			_		_		680
						gaa Glu									-	728
						aac Asn										776
						ctc Leu 250										824
	_					ggg Gly			-		_		-			872
						acc Thr										920
						att Ile										968
						tta Leu										1016
						gcg Ala 330										1064

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Ser Asn Pro Lys Val Lys Leu Tyr Leu Lys Arg Glu Asp Leu Ile His 65 70 75 80	
Gly Gly Ala His Lys Thr Asn Gln Ala Leu Gly Gln Ala Leu Leu Ala 85 90 95	,
Lys Lys Met Gly Lys Thr Arg Ile Ile Ala Glu Thr Gly Ala Gly Gln 100 105 110	
His Gly Val Ala Thr Ala Ile Ala Cys Ala Leu Leu Asn Leu Lys Cys 115 120 125	
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Phe Arg Met His Leu Leu Gly Ala Glu Val Arg Glu Val Asn Ser Gly 145 150 155 160	
Ser Ala Thr Leu Lys Asp Ala Val Asn Glu Ala Leu Arg Asp Trp Ala 165 170 175	
Ser Ser Tyr Lys Asp Thr His Tyr Leu Leu Gly Thr Ala Ala Gly Pro 180 185 190	
His Pro Tyr Pro Thr Met Val Lys Thr Phe Gln Lys Met Ile Gly Asp 195 200 205	
Glu Val Lys Ser Gln Ile Leu Glu Lys Glu Asn Arg Leu Pro Asp Tyr 210 215 220	
Val Ile Ala Cys Val Gly Gly Ser Asn Ala Ile Gly Ile Phe Ser 225 230 235 240	
Ala Phe Leu Asn Asp Lys Glu Val Lys Leu Ile Gly Val Glu Pro Ala	

245 250 Gly Leu Gly Leu Glu Thr Asn Lys His Gly Ala Thr Leu Asn Lys Gly 265 Arg Val Gly Ile Leu His Gly Asn Lys Thr Tyr Leu Leu Gln Asp Asp 275 280 Glu Gly Gln Ile Ala Glu Ser His Ser Ile Ser Ala Gly Leu Asp Tyr 295 Pro Gly Val Gly Pro Glu His Ser Tyr Leu Lys Glu Ser Gly Arg Ala 310 315 Val Tyr Glu Ser Ala Ser Asp Ala Glu Ala Leu Glu Ala Phe Lys Leu 325 330 Leu Cys Gln Lys Glu Gly Ile Ile Pro Ala Leu Glu Ser Ser His Ala 340 345 Leu Ala Tyr Ala Leu Lys Leu Ala Gln Lys Cys Glu Glu Ser Ile 355 360 Ile Val Val Asn Leu Ser Gly Arg Gly Asp Lys Asp Leu Ser Thr Val 375 Tyr Asn Ala Leu Lys Gly Gly Leu Lys 390 <210> 173 <211> 559 <212> DNA <213> Helicobacter pylori <220> <221> CDS <222> (31)...(513) <400> 173 caggtagctt tgggcgagaa aggagagagc atg aat gtc aaa aat cgt ttg agc 54 Met Asn Val Lys Asn Arg Leu Ser gat tgg gaa tat caa tgg gca gtg gct cta gtc tat acg ata tgt atc 102 Asp Trp Glu Tyr Gln Trp Ala Val Ala Leu Val Tyr Thr Ile Cys Ile 10 tcc ata aac gct agg att ttt tat gac ata gat ggt tca gct agc gat 150 Ser Ile Asn Ala Arg Ile Phe Tyr Asp Ile Asp Gly Ser Ala Ser Asp tcg att ttt gac cct aaa aat agc tat tat atg tgg cta gtg ggt cta 198 Ser Ile Phe Asp Pro Lys Asn Ser Tyr Tyr Met Trp Leu Val Gly Leu 55 ata gcg gct ttg ttg tct aac ctt tta ttt gac cca cga ggt agg gat 246 Ile Ala Ala Leu Leu Ser Asn Leu Leu Phe Asp Pro Arg Gly Arg Asp 60 65 tgt tat aaa tct ttc caa gta aga tac cct agg ttt ctc aaa gcc att 294 Cys Tyr Lys Ser Phe Gln Val Arg Tyr Pro Arg Phe Leu Lys Ala Ile 75

Arg Leu Arg Asp Phe Tyr Val Met Leu Leu Thr Ile Pro Phe Ile Ala 110 110 115 120  gct atc cat gag gtt tcg gcg tat tac ggg cat cct agc aac ttc ctt Ala Ile His Glu Val Ser Ala Tyr Tyr Gly His Pro Ser Asn Phe Leu 125 130 135  ata gag ggt ttg gtc att ctt ggc ctt gtg tgt ttt ggg att tgt 48  Ile Glu Gly Leu Val Ile Leu Gly Leu Val Cys Val Phe Gly Ile Cys 140 145 150  tct agg ctt tgc gct aaa tta ggg tgg tgatttaact caaatagcat 53  Ser Arg Leu Cys Ala Lys Leu Gly Trp 155	Phe												gtg Val				342
Ala Ile His Glu Val Ser Ala Tyr Tyr Gly His Pro Ser Asn Phe Leu 135  ata gag ggt ttg gtc att ctt ggc ctt gtg tgt gtt ttt ggg att tgt 48 Ile Glu Gly Leu Val Ile Leu Gly Leu Val Cys Val Phe Gly Ile Cys 140  tct agg ctt tgc gct aaa tta ggg tgg tgatttaact caaatagcat 53 Ser Arg Leu Cys Ala Lys Leu Gly Trp 160  taaatggagg ggggagtaaa aaatta 55  <210> 174 <211> 161 <212> PRT <213> Helicobacter pylori  <400> 174 Met Asn Val Lys Asn Arg Leu Ser Asp Trp Glu Tyr Gln Trp Ala Val 1 1 5 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	Arg					Tyr					Thr					Ala	390
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35 40 45	Asp	Ile		Gly	Ser	Ala	Ser		Ser	Ile	Phe	Asp		Lys	Asn	Ser	
Tyr Tyr Met Trp Leu Val Gly Leu Ile Ala Ala Leu Leu Ser Asn Leu 50 55 60	Tyr		Met	Trp	Leu	Val		Leu	Ile	Ala	Ala		Leu	Ser	Asn	Leu	
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Phe Tyr Asn Ala Val Leu Gly Ser Arg Leu Arg Asp Phe Tyr Val Met 100 105 110	Phe	Tyr	Asn		Val	Leu	Gly	Ser		Leu	Arg	Asp	Phe		Val	Met	
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170

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Val Gln Leu Phe Thr Thr Ile Leu Tyr Leu Phe Ile Gly Leu Tyr Tyr
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                                        205
                                                                   732
ttt aac cgg tgg aat aag tca tgc aag cag tgattttagc gaatggggag
Phe Asn Arg Trp Asn Lys Ser Cys Lys Gln
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gatggggctg ttacctca
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Phe Ile Asn Leu Leu Ala Gly Leu Ser Gly Val Leu Tyr Ala Phe Phe
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Ala Gly Glu Arg Gln Thr Ile Cys Phe Val Phe Gly Leu Val Tyr Asn
                        55
                                            60
Leu Ser Tyr Ala Tyr Val Ala Tyr Gln Trp Lys Leu Asn Ala Asp Val
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Ile Leu Cys Leu Phe Leu Tyr Met Pro Val Thr Ile Tyr Gly Leu Phe
                85
                                    90
Ala Trp Lys Lys Thr Glu Gln His Glu Gly Val Ile Lys Ala Gln Lys
                                105
Leu Ser Lys Asn Trp Arg Phe Ile Leu Ile Leu Gly Val Gly Val Leu
        115
                            120
                                                125
Thr Cys Val Ser Ala Leu Phe Phe Lys Glu Ile Lys Thr Asn Phe Leu
                        135
                                            140
Trp Ala Glu Ser Phe Asn Phe Val Ile Phe Ile Ile Ala Phe Ile Leu
                   150
                                        155
Gln Val Leu Arg Tyr Ile Glu Asn Tyr Ala Leu Val Thr Leu Gly Asn
                165
                                    170
Ile Val Ser Ile Ile Val Trp Phe Cys Ile Phe Gln Ile Ser Thr Glu
                                185
Ser Leu Val Gln Leu Phe Thr Thr Ile Leu Tyr Leu Phe Ile Gly Leu
        195
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Tyr Tyr Phe Asn Arg Trp Asn Lys Ser Cys Lys Gln
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<213> Helicobacter pylori

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cac aaa cac gga tac cgc aag ctt ggg aga acc agc tcg cac aga aag
                                                                   104
His Lys His Gly Tyr Arg Lys Leu Gly Arg Thr Ser Ser His Arg Lys
gcg tta tta aag aat tta gcg atc gct ttg att gag cat aac aaa att
                                                                   152
Ala Leu Leu Lys Asn Leu Ala Ile Ala Leu Ile Glu His Asn Lys Ile
                         25
                                                                   200
gaa aca ggg att tat aag gct aag gaa ttg cgc agt tac att gag aaa
Glu Thr Gly Ile Tyr Lys Ala Lys Glo Leo Arg Ser Tyr Ile Glu Lys
                                                                   248
ttg acg aca gcg gct cgt gtg ggc gat ttt aat gcg cac cgc cat gtt
Leu Thr Thr Ala Ala Arg Val Gly Asp Phe Asn Ala His Arg His Val
ttt gca tat ttg caa aac aaa gaa gcc acc cac aag ctt gta act gaa
                                                                   296
Phe Ala Tyr Leu Gln Asn Lys Glu Ala Thr His Lys Leu Val Thr Glu
atc gcg ccc aaa tac gcg caa agg aat ggc gga tac acc agg atc caa
                                                                   344
Ile Ala Pro Lys Tyr Ala Gln Arg Asn Gly Gly Tyr Thr Arg Ile Gln
         85
                             90
cgc acc act ttt aga aga ggg gac gct tcc act cta gcc acc att gaa
                                                                   392
Arg Thr Thr Phe Arg Arg Gly Asp Ala Ser Thr Leu Ala Thr Ile Glu
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ttt gta tgaaatttga tgaactgcta gccaagattt agtcttggtt ggtggttatc
                                                                   448
Phe Val
115
                                                                   451
gct
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Arg Lys Ala Leu Leu Lys Asn Leu Ala Ile Ala Leu Ile Glu His Asn
Lys Ile Glu Thr Gly Ile Tyr Lys Ala Lys Glu Leu Arg Ser Tyr Ile
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Glu Lys Leu Thr Thr Ala Ala Arg Val Gly Asp Phe Asn Ala His Arg His Val Phe Ala Tyr Leu Gln Asn Lys Glu Ala Thr His Lys Leu Val 70 75 Thr Glu Ile Ala Pro Lys Tyr Ala Gln Arg Asn Gly Gly Tyr Thr Arg 90 Ile Gln Arg Thr Thr Phe Arg Arg Gly Asp Ala Ser Thr Leu Ala Thr 105 100 Ile Glu Phe Val 115 <210> 179 <211> 1204 <212> DNA <213> Helicobacter pylori <220> <221> CDS <222> (36) ... (1142) <400> 179 aaaataagcg ttttgatgcc atttttggag cgatc gtg gaa ttg agt tat tat 53 Val Glu Leu Ser Tyr Tyr gaa att tta gaa gtg gaa aaa cac agc aac caa gag acc att aaa aag 101 Glu Ile Leu Glu Val Glu Lys His Ser Asn Gln Glu Thr Ile Lys Lys 10 15 tct tac aga aag ctg gct tta aaa tac cac cca gac aga aac gcc ggc 149 Ser Tyr Arg Lys Leu Ala Leu Lys Tyr His Pro Asp Arg Asn Ala Gly 25 30 35 gat aaa gaa gcc gaa gaa aaa ttc aag ctc atc aat gaa gcc tat ggg 197 Asp Lys Glu Ala Glu Glu Lys Phe Lys Leu Ile Asn Glu Ala Tyr Gly 40 45 gtg tta agc gat gaa aag aag cgg gcc tta tac gac agg tat ggt aaa 245 Val Leu Ser Asp Glu Lys Lys Arg Ala Leu Tyr Asp Arg Tyr Gly Lys 55 aaa ggc tta aac caa gcc ggc gca agc caa ggc gat ttt tct gat ttt 293 Lys Gly Leu Asn Gln Ala Gly Ala Ser Gln Gly Asp Phe Ser Asp Phe 75 85 ttt gaa gat tta ggc tcg ttt ttt gaa gac gct ttt ggg ttt ggc gct 341 Phe Glu Asp Leu Gly Ser Phe Phe Glu Asp Ala Phe Gly Phe Gly Ala 90 95 agg ggg agt aaa agg caa aaa agc tct atc gca ccg gat tat ttg caa 389 Arg Gly Ser Lys Arg Gln Lys Ser Ser Ile Ala Pro Asp Tyr Leu Gln 110 acc ctt gaa ttg agt ttc aaa gaa gcg gtt ttt ggc tgt aaa aaa acc 437

Thr	Leu 120	Glu	Leu	Ser	Phe	Lys 125	Glu	Ala	Val	Phe	Gly 130	Cys	Lys	Lys	Thr	
		-			cag Gln 140	-		_	_	_						485
-		_		-	cta Leu			_	_		_					533
_			_	_	caa Gln			_	_					_		581
	_			_	ggc Gly	_		_				_			-	629
_					atc Ile			_	_	_		_				677
					gat Asp 220											725
					gga Gly											773
-			-		cat His		-	_	_		-	~				821
					ttc Phe											869
					Gly ggg											917
					act Thr 300											965
					aga Arg											1013
					aat Asn											1061

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His Ala Ser Phe Gly Tyr Glu Gly Glu Pro His Lys Ser Val Leu Glu
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acc tgt att tct aaa att aaa gac tgg ttc aaa taaaaggttg ttgatgcatg 1162
Thr Cys Ile Ser Lys Ile Lys Asp Trp Phe Lys
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Pro Asp Arg Asn Ala Gly Asp Lys Glu Ala Glu Glu Lys Phe Lys Leu
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Ile Asn Glu Ala Tyr Gly Val Leu Ser Asp Glu Lys Lys Arg Ala Leu
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Tyr Asp Arg Tyr Gly Lys Lys Gly Leu Asn Gln Ala Gly Ala Ser Gln
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Gly Asp Phe Ser Asp Phe Phe Glu Asp Leu Gly Ser Phe Phe Glu Asp
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Ala Phe Gly Phe Gly Ala Arg Gly Ser Lys Arg Gln Lys Ser Ser Ile
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Ala Pro Asp Tyr Leu Gln Thr Leu Glu Leu Ser Phe Lys Glu Ala Val
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                                                125
Phe Gly Cys Lys Lys Thr Ile Lys Val Gln Tyr Gln Ser Val Cys Glu
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Ser Cys Asp Gly Thr Gly Ala Lys Asp Lys Ala Leu Glu Thr Cys Lys
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Gln Cys Asn Gly Gln Gly Gln Val Phe Met Arg Gln Gly Phe Met Ser
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                                    170
Phe Ala Gln Thr Cys Gly Ala Cys Gln Gly Lys Gly Lys Ile Val Lys
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                                185
Thr Pro Cys Gln Ala Cys Lys Gly Lys Thr Tyr Ile Leu Lys Asp Glu
                            200
                                                205
Glu Ile Asp Ala Ile Ile Pro Glu Gly Ile Asp Asp Gln Asn Arg Met
                        215
Val Leu Lys Asn Lys Gly Asn Glu Tyr Glu Lys Gly Lys Arg Gly Asp
                    230
                                        235
Leu Tyr Leu Glu Ala Gln Val Lys Glu Asp Glu His Phe Lys Arg Glu
                245
                                    250
Gly Cys Asp Leu Phe Ile Lys Ala Pro Val Phe Phe Thr Thr Ile Ala
                                265
                                                    270
Leu Gly His Thr Ile Lys Val Pro Ser Leu Lys Gly Asp Glu Leu Glu
        275
                            280
Leu Lys Ile Pro Arg Asn Ala Arg Asp Lys Gln Thr Phe Ala Phe Arg
                        295
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Asn Glu Gly Val Lys His Pro Glu Ser Ser Tyr Arg Gly Ser Leu Ile

305					310					313					320	
				325				Lys	330					335		
Glu	Leu	Leu	Glu 340	Lys	Leu	His	Ala	Ser 345	Phe	Gly	Tyr	Glu	Gly 350	Glu	Pro	
His	Lys	Ser 355	Val	Leu	Glu	Thr	Cys 360	Ile	Ser	Lys	Ile	Lys 365	Asp	Trp	Phe	
Lys																
	)> 18															
<212	1> 81 2> Di	ΝA	1		3											
		STIC	obaci	ter p	oylo	rı										
	l> CI			70)												
	2> (: 2- 10		••(7	/9)												
	)> 18 atgga														a ttc	50
			ме і 1	L Arg	g va.	т ме	5 5	я гу:	5 116	e Gil	л ге	1 Let		a r.y:	s Phe	
								cac His								98
1111	15	110	AIU	пси	110	20	DCI		110	Dea	25	цуз	БУБ	vai	Бей	
								cag Gln								146
30	-4-		-4	-1 -	35				-7-	40				202	45	
								tgc Cys								194
				50	_			-	55	-	-		-	60		
								tgc Cys								242
			65					70					75			
								caa Gln								290
		80					85					90				
	Lys					Asn		ctt Leu			Lys					338
	95					100					105					
Val					Ala			gcg Ala		Phe					Pro	386
110					115					120					125	404
								aaa Lys								434

	130	135	140	
	Glu Leu His		gat tca gag cgt ttt tgg Asp Ser Glu Arg Phe Trp 155	482
_	-		atc gtc cat aac gca tgc Ile Val His Asn Ala Cys 170	530
			aag gac agc tac acc act Lys Asp Ser Tyr Thr Thr 185	578
		Glu Cys Tyr I	aaa acg cta ggg ttt atc Lys Thr Leu Gly Phe Ile 200 205	626
			ogo tat tta gaa ttg got Arg Tyr Leu Glu Leu Ala 220	674
	Pro Asp Leu		aaa gaa aaa aag gct ctt Lys Glu Lys Lys Ala Leu 235	722
			gcg tat cgc cac att aaa Ala Tyr Arg His Ile Lys 250	770
aaa cac gcc taa Lys His Ala 255	accccta aact	agccct aatcgcg	gcta g	810
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Ala Leu Pro Asn 20	Ser His Pro	Leu Leu Lys L 25	ys Val Leu Asn Tyr Ala 30	
Lys Lys His Phe 35	Ser Gln Cys	His Met Leu S	Ser Ser Leu Leu Ile 45	
Leu Asn Asp Thr 50	Glu Cys Phe 55	Lys Lys Asn T	Tyr Leu Leu Asn Trp Val 60	
65	70	7	sp Ile Ser Ala His Ser 5 80	
	Leu Gln Lys 85	Ser His Leu P 90	ro Ile Arg Ile Lys Ile 95	
Met Ala Gln Asn	-			

Phe Gly Ala Glu Tyr Ala Leu Phe Ile Thr Lys His Pro Ile Ala Lys 115 120 125 Arg Phe Leu Arg Gln Lys Phe Ser Gly Cys Val Phe Leu Glu Thr Gln 135 140 Asp Glu Leu His Ile Arg Gly Asp Ser Glu Arg Phe Trp Glu Leu Ile 150 Val Thr Leu Asn Glu Asn Arg Ile Val His Asn Ala Cys Leu Asp Phe 165 170 Ile Tyr Pro Asn Gly Phe Gly Lys Asp Ser Tyr Thr Thr Met Ala Glu 185 190 Arg Lys Leu Lys Glu Cys Tyr Lys Thr Leu Gly Phe Ile Lys His Glu 200 205 Asp Phe Ser Glu Val Lys Lys Arg Tyr Leu Glu Leu Ala Lys Thr Tyr 215 220 His Pro Asp Leu Cys Asp Leu Lys Glu Lys Lys Ala Leu Tyr Ala Lys 230 235 Arg Phe Ala Ile Ile Gln Glu Ala Tyr Arg His Ile Lys Lys His Ala 245 250 <210> 183 <211> 900 <212> DNA <213> Helicobacter pylori <220> <221> CDS <222> (59)...(778) <400> 183 catgatgatt attataaaat cctaacgccg catgaacaaa taggatgggt caaaaaag 58 atg aag tca aat aaa aag tcc aat cgt tta aga gcg att tat aga gct 106 Met Lys Ser Asn Lys Lys Ser Asn Arg Leu Arg Ala Ile Tyr Arg Ala 1 tta gtg atc gct ata gga cta gct gtt atc atc gtt ttc aat tac ttt 154 Leu Val Ile Ala Ile Gly Leu Ala Val Ile Ile Val Phe Asn Tyr Phe 20 aac cgc aaa aac aat aac gcc cgc tcc agc cgt agg gct tgt tcg tgc 202 Asn Arg Lys Asn Asn Asn Ala Arg Ser Ser Arg Arg Ala Cys Ser Cys 35 ttt ttt tcc ctt acc ggg gtt aat tta gaa aaa ata ggc act ttt gat 250 Phe Phe Ser Leu Thr Gly Val Asn Leu Glu Lys Ile Gly Thr Phe Asp 50 acg gac gct aaa ctc att gtc tta aac cac caa agc tta cta gac atc 298 Thr Asp Ala Lys Leu Ile Val Leu Asn His Gln Ser Leu Leu Asp Ile 70 75 att tat tta gaa gcc tac cac cct aga aat att tgc tgg atc gct aaa 346 Ile Tyr Leu Glu Ala Tyr His Pro Arg Asn Ile Cys Trp Ile Ala Lys

aaa gag ctg Lys Glu Leu							394
gga atg att Gly Met Ile 115							442
ttg aaa gcg Leu Lys Ala 130					Pro Leu		490
ttc cct gaa Phe Pro Glu 145		g Gly Lys					538
aag caa ggg Lys Gln Gly							586
ccc atg gtg Pro Met Val					•		634
gaa gcc tat Glu Ala Tyr 195							682
cct gat ttt Pro Asp Phe 210	•		_	-	Gln Glu		730
caa aaa gag Gln Lys Glu 225		s His Tyr	_		_	_	778
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Leu Val Ile	20		25		30		
Asn Arg Lys		40			45	_	
Phe Phe Ser 50		55		60			
Thr Asp Ala	Lys Leu Il	e Val Leu	Asn His	Gln Ser	Leu Leu	Asp Ile	

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Ile Tyr Leu Glu Ala Tyr His Pro Arg Asn Ile Cys Trp Ile Ala Lys
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Lys Glu Leu Gly Glu Ile Pro Phe Tyr Gly His Ala Leu Thr Asp Thr
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                                105
Gly Met Ile Leu Ile Asp Arg Glu Asp Lys Lys Gly Ile Val Ser Leu
                            120
Leu Lys Ala Cys Lys Glu Lys Leu Asp Gln Asn Arg Pro Leu Val Ile
                        135
                                            140
Phe Pro Glu Gly Thr Arg Gly Lys Gly Glu Lys Phe Leu Pro Phe
                    150
                                        155
Lys Gln Gly Ala Lys Ile Ile Ala Glu Lys Phe Gln Leu Lys Ile Gln
                165
                                    170
Pro Met Val Leu Ile Asn Ser Ile Lys Ile Phe Asn Ser Lys Pro Leu
                                185
Glu Ala Tyr Lys Ala Arg Thr Arg Leu Val Met Leu Glu Ser Tyr Thr
                            200
Pro Asp Phe Asn Ser Pro Thr Trp Tyr Glu Glu Leu Gln Glu Arg Met
                        215
                                            220
Gln Lys Glu Tyr Leu Lys His Tyr His Glu Leu Asn Pro Ser Glu Gln
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                                           Met Glu Ile Arg Thr
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                                                                   102
Phe Leu Glu Arg Ala Leu Lys Glu Asp Leu Gly His Gly Asp Leu Phe
gaa agg gtg tta gaa aaa gat ttt aag gcc aca gcg ttt gtt agg gct
                                                                   150
Glu Arg Val Leu Glu Lys Asp Phe Lys Ala Thr Ala Phe Val Arg Ala
             25
                                 30
aaa caa gag ggc gtg ttt tca ggc gaa aaa tac gct tta gag ttg ttg
                                                                   198
Lys Gln Glu Gly Val Phe Ser Gly Glu Lys Tyr Ala Leu Glu Leu Leu
         40
                             45
gaa atg acc ggc att gaa tgc gtt caa acg att aag gat aaa gaa cgc
                                                                   246
Glu Met Thr Gly Ile Glu Cys Val Gln Thr Ile Lys Asp Lys Glu Arg
    55
                         60
ttc aag cct aaa gac gct tta atg gaa att agg ggg gat ttt agc atg
                                                                   294
Phe Lys Pro Lys Asp Ala Leu Met Glu Ile Arg Gly Asp Phe Ser Met
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70

70				75				80					85	
	_	•		_		ctt Leu			_			_	_	342
						cgt Arg								390
	 _	_	_	_	_	aga Arg 125	_	_						438
						aat Asn								486
					_	ctt Leu	_	_						534
						acg Thr								582
						tgc Cys								630
_						att Ile 205			-		_	-	_	678
						gct Ala								726
<b>-</b>		7. 4		_		aac Asn				_			_	774
						gcc Ala								822
						cac His				taaa	igact	tt		868
													gcctg	

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                        55
                                            60
Lys Asp Lys Glu Arg Phe Lys Pro Lys Asp Ala Leu Met Glu Ile Arg
                                        75
Gly Asp Phe Ser Met Leu Leu Lys Val Glu Arg Thr Leu Leu Asn Leu
                                    90
Leu Gln His Ser Ser Gly Ile Ala Thr Leu Thr Ser Arg Phe Val Glu
            100
                                105
Ala Leu Asn Ser His Lys Val Arg Leu Leu Asp Thr Arg Lys Thr Arg
                            120
Pro Leu Leu Arg Ile Phe Glu Lys Tyr Ser Val Leu Asn Gly Gly Ala
                        135
                                            140
Ser Asn His Arg Leu Gly Leu Asp Asp Ala Leu Met Leu Lys Asp Thr
                    150
                                        155
His Leu Arg His Val Lys Asp Leu Lys Ser Phe Leu Thr His Ala Arg
                165
                                    170
Lys Asn Leu Pro Phe Thr Ala Lys Ile Glu Ile Glu Cys Glu Ser Phe
                                185
Glu Glu Ala Lys Asn Ala Met Asn Ala Gly Ala Asp Ile Val Met Cys
                            200
Asp Asn Leu Ser Val Leu Glu Thr Lys Glu Ile Ala Ala Tyr Arg Asp
                        215
                                            220
Ala His Tyr Pro Phe Val Leu Leu Glu Ala Ser Gly Asn Ile Ser Leu
                    230
                                        235
Glu Ser Ile Asn Ala Tyr Ala Lys Ser Gly Val Asp Ala Ile Ser Val
                245
                                    250
Gly Ala Leu Ile His Gln Ala Thr Phe Ile Asp Met His Met Lys Met
            260
Ala
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<211> 1153
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                                                        Met Ile
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Met Glu Ile Arg Thr Phe Leu Glu Arg Ala Leu Lys Glu Asp Leu Gly

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<400> 186

<213> Helicobacter pylori

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	_						-		_	-	tgg Trp 30		-	_	_	152
	_	_		_			_			_	atc Ile		_			200
_	-		_	_	_	_		_	_		atg Met		-		_	248
		-				-	-			-	gag Glu		_	_		296
_						_	_	_	_		tta Leu		_			344
											tcc Ser 110					392
											att Ile					440
											cgt Arg					488
											gag Glu					536
											cat His					584
											gtg Val 190					632
											gta Val					680
tgc	cct	tta	ttg	aat	tta	aag	ctt	tgc	act	ctg	gat	tta	tcc	aaa	cga	728

Cys Pro Leu Le	eu Asn Leu Lys 215	Leu Cys Thr 220	Leu Asp Leu Ser	Lys Arg 225
•	r Met Arg Ty		ttt tta aaa gag Phe Leu Lys Glu 240	
			tat gcg gac gct Tyr Ala Asp Ala 255	
_	-	Cys Leu Leu	aac ttg ccc agc Asn Leu Pro Ser 270	
			aaa tgt gct aac Lys Cys Ala Asn 285	
	_	-	gag att cca tcg Glu Ile Pro Ser	
	u Lys Thr Lys		aat cat aaa agc Asn His Lys Ser 320	-
			ggg act gaa att Gly Thr Glu Ile 335	
tct cta ttt ga Ser Leu Phe As 340		Cys Cys Leu	gta ttt tgatttt Val Phe 350	atc 1110
ggttacttcg cac	tcatcgt atato	ttttt gtattct	tgt atg	1153
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20		25	30 His Lys Asp Ile	
35		40	45 Ser Leu Gly Met	
50	55		60	
65	70		Ile Asn Leu Glu 75	80
ьуѕ Ala Cys Gl	y Leu Asn Phe 85	Ile Glu Thr 90	Ser Glu Asn Leu	Ile Thr 95

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Leu Lys Asn Leu Lys Thr Pro Leu Lys Asp Glu Val Phe Ser Phe Ile
                                105
Asp Leu Glu Thr Thr Gly Ser Cys Pro Ile Lys His Glu Ile Leu Glu
        115
                            120
                                                125
Ile Gly Ala Val Gln Val Lys Gly Glu Ile Ile Asn Arg Phe Glu
                       135
Thr Leu Val Lys Val Lys Ser Val Pro Asp Tyr Ile Ala Glu Leu Thr
                    150
                                        155
Gly Ile Thr Tyr Glu Asp Thr Leu Asn Ala Pro Ser Ala His Glu Ala
                165
                                    170
Leu Gln Glu Leu Arg Leu Phe Leu Gly Asn Ser Val Phe Val Ala His
                                185
Asn Ala Asn Phe Asp Tyr Asn Phe Leu Gly Arg Tyr Phe Val Glu Lys
                            200
Leu His Cys Pro Leu Leu Asn Leu Lys Leu Cys Thr Leu Asp Leu Ser
Lys Arg Ala Ile Leu Ser Met Arg Tyr Ser Leu Ser Phe Leu Lys Glu
                    230
                                        235
Leu Leu Gly Phe Gly Ile Glu Val Ser His Arg Ala Tyr Ala Asp Ala
                245
                                    250
Leu Ala Ser Tyr Lys Leu Phe Glu Ile Cys Leu Leu Asn Leu Pro Ser
            260
                                265
Tyr Ile Lys Thr Thr Met Asp Leu Ile Asp Phe Ser Lys Cys Ala Asn
                            280
                                                285
Thr Leu Ile Lys Arg Pro Pro Lys Ala Arg Tyr Gln Glu Ile Pro Ser
                        295
Pro Phe Ser Leu Phe Glu Lys Thr Lys Gly Leu Phe Asn His Lys Ser
                    310
                                        315
Asn Gln Leu Asn Glu Ser Cys Leu Met Gly Phe Met Gly Thr Glu Ile
                325
Leu Ala Ser Leu Phe Asp Thr Phe Glu Cys Cys Leu Val Phe
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                                                                  105
Lys Ile Glu Ser Asn Asp Ser His Leu Arg Gly Ile Leu Lys Asp Glu
                             10
ctc tac tat caa atc ccc atc tac caa cgc cct tat caa tgg aca gaa
                                                                  153
Leu Tyr Tyr Gln Ile Pro Ile Tyr Gln Arg Pro Tyr Gln Trp Thr Glu
                         25
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							gac Asp									201
							tgc Cys									249
_		_			-		acc Thr		_	_	_	_		_		297
_		_				-	ctt Leu 90	_		-		_	_			345
	-	_		-		-	aat Asn									393
	-				-	_	aaa Lys	_	_	-						441
		_	_		_		caa Gln	_	_		_				_	489
							aat Asn									537
		-			-		ctc Leu 170	_								585
	_					_	tat Tyr			_						633
							gca Ala									681
							gcg Ala									729
							gag Glu									777
							tcg Ser									825

245 250 255

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Ala Gly Gly Arg Glu Val Val Phe Lys Asp Asn Asp Lys Lys Glu Ala
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Lys Leu Phe Glu Tyr Gly Ile Asn Ala Val Val Leu Gly Asp Tyr Leu
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		-			aag Lys									_		152
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Ala	Gly	Leu	20 Gly	Gly	Glu	Ser	Val	25 Thr	Pro	Asn	Val	Leu	30 Lys	Asp	Phe	
Gly		35 Met	Leu	Ala	Gln		40 Val	Gln	Phe	Gln		45 Tyr	Gly	Ser	Leu	
Gly 65	50 Leu	Arg	Met	Gly	Asp 70	55 Lys	His	His	Thr		60 Glu	Leu	Ser	Thr		
	His	Gly	Asp	Ala 85	Pro	Ser	Cys	Ser	Leu 90	75 Lys	Lys	Leu	Lys	Ser 95	80 Cys	
Glu	Ser	Ala	Ara		T.011	Gln	α1 ۵	Laze		Dro	λνα	C111	т1 о		C1.,	

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		aac Asn 5				-			-		_	_	_			106
		gat Asp														154
		cgt Arg														202
		gcg Ala														250
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	Glu	gaa Glu 85	Leu	Asp		Asn	Lys	Val					Glu			346
		aat Asn														394
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		gcg Ala 165														586
		ggc Gly		_				_	-				-			634
•		gac Asp										_		_	_	682
		ggg														730
		acg Thr														778
		ttg Leu 245														826
		att Ile														874
		atc Ile												-		922
		atc Ile														970
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		ctt Leu 325														1066
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		caa Gln														1162

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		tat Tyr 405														1306
_	_	aaa Lys		_	_		-				_	_				1354
		gaa Glu														1402
		att Ile												_		1450
		cct Pro										-			_	1498
		aaa Lys 485														1546
		ggg Gly														1594
		cct Pro														1642
		cac His														1690
		tta Leu														1738
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caa Gln	agc Ser	aat Asn	ttg Leu	agt Ser	att Ile	gat Asp	aca Thr	agc Ser	atc Ile	gtg Val	caa Gln	caa Gln	aga Arg	gaa Glu	gaa Glu	1834

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			_	-	_								ggg Gly 640			1978
													aac Asn			2026
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Pro His Tyr Lys Met Asn Leu Asn Ser Lys Ala Ser Val Lys Ile Thr

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Glu His Arg His Ile Ser Ile Lys Pro Asn Asp Leu Val Ile Ile Ser
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Phe Leu Ile Lys Lys Glu Ala Lys Val Ala Tyr Gln Glu Phe Asp Asn
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Ile His Val Ser Gly His Ala Ala Gln Glu Glu Gln Lys Leu Met Leu
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His Val Ala Arg His Lys Gln Thr Ala Ile Ser Cys Gly Val Pro Glu
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Lys Asn Ile Tyr Leu Met Glu Asp Gly Asp Gln Val Glu Val Gly Pro
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Ala Phe Ile Lys Lys Val Gly Thr Ile Lys Ser Gly Lys Ser Tyr Val
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Lys Asn Lys Gln Ala Leu Leu Glu Ser Ser Gln Phe Ser Ser Leu Gly

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			_	_	_						gtt Val		-			489
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				_					_		aac Asn				_	585
											acc Thr		-			633
							-			-	gca Ala	_				681
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											agg Arg 240					777
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											ccg Pro					873
											tta Leu					921
											aaa Lys					969
Lys	Ser 310	Gln	Glu	Tyr	Leu	Ala 315	Lys	Ala	Leu	Gly	caa Gln 320	Asp	Tyr	Gln	Glu	1017
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		_	_		aga Arg		-				-				_	1833
_			_		cct Pro					_						1881
-					tta Leu		_									1929
-		_			tta Leu		_				-		_			1977
	_	-			ttt Phe 650				-			-		_		2025
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	_				atc Ile						_					2169
					tac Tyr											2217
					aat Asn 730											2265
					gct Ala											2313
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atg cgg tta Met Arg Leu		_	_	_	_	
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ctt tta agg Leu Leu Arg						
att tcc aac Ile Ser Asn						
cct tat ttt Pro Tyr Phe 145	Tyr Ala P		Asn Gln			
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gac ttg ctc Asp Leu Leu 175	Val Leu A					ie
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ccc gca ttc Pro Ala Phe						

Val										gtg Val						783
										ccc Pro						831
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										gac Asp						927
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tcg ctt gat Ser Leu Asp							
cct tat ttt Pro Tyr Phe							
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Pro Pro Leu 35		4	10		45		
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Lys Tyr Ala Gln Thr Phe His Ser Asn Pro Tyr Thr Asn Gln Lys Thr
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Pro Asn Ser Asp Leu Tyr Tyr Pro Ala Leu Asn Glu Gly Asn Ser Phe
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Ser Ile Gln Ile Met Gly Ile Ser Val Ala Glu Leu Leu Lys Ser Lys
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Lys Phe Leu Ser Leu Asp Val Ser Phe Lys Lys Gly Ser Val Leu Trp
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Gly Gly Arg Pro Tyr Phe Ser Glu Val Gly Glu Phe Met Gly Met Ala
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Asp Trp His Ile Gln Lys Val Glu Met Asn His Gln Val Tyr Asp Ile
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Glu Thr Met Leu Ala Asp Ser Ala Phe Arg Glu His Glu Glu Glu Gln
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                                    35
                                                       40
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					tgg Trp											450
					aaa Lys											498
					gag Glu											546
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Glu 65	Lys	Arg	Lys	Arg	Trp 70	Tyr	Glu	Leu	Phe	Lys 75	Lys	Lys	Pro	Lys	Pro 80	
Lys	Ser	Ser	Met	Gly 85	Glu	Phe	Val	Phe	Asp 90	Gln	Lys	Glu	Asn	Arg 95	Ile	
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Glu Ser Tyr Gly Asp Lys Asn Leu Phe Thr Leu Ile Ser Ile Ser Pro
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Val Asn Tyr Gln Val Ser Met Ile Gln Asn Gln Thr Lys Met Val Asn
gac aac ggc ttg caa aag cct ttg ata aag ttt ccg cct tac gca gga
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Asp Asn Gly Leu Gln Lys Pro Leu Ile Lys Phe Pro Pro Tyr Ala Gly
gcg ggt ttt gaa gtg ggc tat aag caa ttt ttt ggt aag aaa aaa tgg
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Ala Gly Phe Glu Val Gly Tyr Lys Gln Phe Phe Gly Lys Lys Lys Trp
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Gly Val Met Lys Lys Gly Ile Pro Val Gly Asp Ser Gly Phe Ile Tyr
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Asn Ser Phe Ser Phe Gly Gly Asn Thr Leu Thr Glu Arg Asp Ser Tyr
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Gln Gly Gln Tyr Tyr Val Asn Leu Phe Thr Tyr Gly Val Gly Leu Asp
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Thr Leu Trp Asn Phe Val Asn Lys Glu Asn Met Val Phe Gly Phe Val
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Val Gly Ile Gln Leu Ala Gly Asp Ser Trp Ala Thr Ser Ile Ser Lys
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                                        180
gcc aat ttc cag ttt tta tgg aag ttt ggg gtc cgc acc cat atc gct
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Ala Asn Phe Gln Phe Leu Trp Lys Phe Gly Val Arg Thr His Ile Ala
aaa cac aat agc cta gaa tta ggg att aaa gtg cct acg atc aca cac
                                                                   678
Lys His Asn Ser Leu Glu Leu Gly Ile Lys Val Pro Thr Ile Thr His
cag ctt ttc tct ctt acc aac gaa aag gga tac acc tta cag gct gat
                                                                   726
Gln Leu Phe Ser Leu Thr Asn Glu Lys Gly Tyr Thr Leu Gln Ala Asp
gtg cgt aga gtt tat gcg ttt caa atc agt tac ttg agg gat ttt
                                                                   771
Val Arg Arg Val Tyr Ala Phe Gln Ile Ser Tyr Leu Arg Asp Phe
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Leu Ile Lys Phe Pro Pro Tyr Ala Gly Ala Gly Phe Glu Val Gly Tyr
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Lys Gln Phe Phe Gly Lys Lys Trp Phe Gly Met Arg Tyr Tyr Gly
                    70
                                        75
Phe Phe Asp Tyr Ala His Asn Arg Phe Gly Val Met Lys Lys Gly Ile
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                                    90
Pro Val Gly Asp Ser Gly Phe Ile Tyr Asn Ser Phe Ser Phe Gly Gly
                                105
Asn Thr Leu Thr Glu Arg Asp Ser Tyr Gln Gly Gln Tyr Tyr Val Asn
                            120
                                                125
Leu Phe Thr Tyr Gly Val Gly Leu Asp Thr Leu Trp Asn Phe Val Asn
                        135
Lys Glu Asn Met Val Phe Gly Phe Val Val Gly Ile Gln Leu Ala Gly
                    150
                                        155
Asp Ser Trp Ala Thr Ser Ile Ser Lys Glu Ile Ala His Tyr Ala Lys
                165
His His Ser Asn Ser Ser Tyr Ser Pro Ala Asn Phe Gln Phe Leu Trp
                                185
Lys Phe Gly Val Arg Thr His Ile Ala Lys His Asn Ser Leu Glu Leu
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Gly Ile Lys Val Pro Thr Ile Thr His Gln Leu Phe Ser Leu Thr Asn
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Glu Lys Gly Tyr Thr Leu Gln Ala Asp Val Arg Arg Val Tyr Ala Phe

235

230

Gln Ile Ser Tyr Leu Arg Asp Phe

				aac Asn									663
				aat Asn									711
				agc Ser 180	-			_	_	-		_	759
		-		aat Asn	_	_		_		 -	-		807
				act Thr									855
	-		-	gaa Glu							-		903
	_		_	gat Asp			_		_				951
				aaa Lys 260									999
_		_		gcc Ala				_			_		1047
				agc Ser									1095
				caa Gln									1143
				gat Asp									1191
				att Ile 340									1239
				gag Glu									1287

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Lys Gln Thr Ile Gln Gly Val Ala Gln Tyr Cys Asp Ala Lys Ile Gln
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Ala Thr Lys Glu Val Ile Ser Gln Glu Ala Pro Val Gln Lys Asp Ser
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Leu Phe
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gcaaccegge tgtgccgccg tattcatcaa tcaccaaage ggt atg gat ttg ctc
                                                 Met Asp Leu Leu
ttt ttt cat ttt aat aag gat ttg aga aat gga agc gct ttc ggg gac
                                                                    223
Phe Phe His Phe Asn Lys Asp Leu Arg Asn Gly Ser Ala Phe Gly Asp
gat gat cat tit cct aac gat tig att gaa atc atg cat tit ggg ggt
                                                                    271
Asp Asp His Phe Pro Asn Asp Leu Ile Glu Ile Met His Phe Gly Gly
                                      30
                                                          35
aaa aat aga gcg cga aag caa atc cct aat atg cac cat gcc gat aat
                                                                   319
Lys Asn Arg Ala Arg Lys Gln Ile Pro Asn Met His His Ala Asp Asn
             40
                                 45
                                                      50
gtt atc ctt aga acc ctt gca ata agg gta gcg cgt gaa atg gcc ttt
                                                                   367
Val Ile Leu Arg Thr Leu Ala Ile Arg Val Ala Arg Glu Met Ala Phe
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taaaacaatg totatatttt ottoatagot gttttottoa tooaaacaca coatgtottt 427
tcgtggggtc atgatttctt tag
                                                                   450
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His Phe Gly Gly Lys Asn Arg Ala Arg Lys Gln Ile Pro Asn Met His
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	ıl Thr Leu Va		aaa att gcc ggt Lys Ile Ala Gly 160	
_			aga gcg gtc att Arg Ala Val Ile 175	
	•	s Glu Ile Asp	cct tta gtg cca Pro Leu Val Pro 190	
			acc gag tgc tat Thr Glu Cys Tyr 205	
			acc ctt ttt gca Thr Leu Phe Ala	
	er Asn Ser Le		tac tac att aaa Tyr Tyr Ile Lys 240	
_		-	cta gag gct tta Leu Glu Ala Leu 255	
		Glu Cys Gln	acc cct tta aag Thr Pro Leu Lys 270	
			gtg cac aac att Val His Asn Ile 285	-
			ggc ttg ata gcg Gly Leu Ile Ala	
caa tcc ttt ga Gln Ser Phe Gl 31	u Lys Leu Leu		tgatttttaa gggt1	tttttg 1018
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Tyr Lys Asp Glu Ile Ala Lys Leu Lys Gly Lys Lys Ala Lys Glu Ile
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Tyr Glu Thr Leu Ala Leu Lys Asp Ile Leu Gln Ala Ser Ser Ala Leu
                    70
                                        75
Met Pro Leu Tyr Glu Lys Asp Pro Asn Asn Gly Tyr Ile Ser Leu Glu
                                    90
                85
Ile Asp Pro Phe Leu Glu Asp Asp Ala Ala Lys Ser Ile Asp Glu Ala
            100
                                105
Lys Arg Leu Phe Lys Thr Leu Asn Arg Pro Asn Val Met Ile Lys Val
                            120
Pro Ala Ser Glu Ser Gly Ile Glu Val Val Ser Ala Leu Thr Gln Ala
                        135
Ser Ile Pro Val Asn Val Thr Leu Val Phe Ser Pro Lys Ile Ala Gly
                    150
                                        155
Glu Ile Ala Gln Ile Leu Ala Lys Glu Ala Gln Lys Arg Ala Val Ile
               165
                                    170
Ser Val Phe Val Ser Arg Phe Asp Lys Glu Ile Asp Pro Leu Val Pro
            180
                               185
                                                     190
Lys Asn Leu Gln Ala Gln Ser Gly Ile Ile Asn Ala Thr Glu Cys Tyr
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                            200
                                                205
Tyr Gln Ile Asn Gln His Ala Asn Lys Leu Thr Ser Thr Leu Phe Ala
                        215
                                            220
Ser Thr Gly Val Lys Ser Asn Ser Leu Ala Lys Asp Tyr Tyr Ile Lys
                    230
                                        235
Ala Leu Cys Phe Lys Asn Ser Ile Asn Thr Ala Pro Leu Glu Ala Leu
                245
                                    250
Asn Ala Tyr Leu Leu Asp Pro Asn Thr Glu Cys Gln Thr Pro Leu Lys
                                265
Thr Thr Glu Ile Glu Ala Phe Lys Lys Glu Leu Lys Val His Asn Ile
                            280
Asp Leu Glu Asn Thr Ala Gln Lys Leu Leu Lys Glu Gly Leu Ile Ala
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Phe Lys Gln Ser Phe Glu Lys Leu Leu Ser Ser Phe
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                                                             Met
ctc aat ttt atg aca aag aag aaa aat aga atg caa gat tgc aaa atg
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Glu Asn Asp Phe Leu Lys Leu Ile Asn Lys Gly Ala Ile Cys Gly Ala

Leu	Asn	Phe	Met 5	Thr	Lys	Lys	Lys	Asn 10	Arg	Met	Gln	Asp	Cys 15	Lys	Met	
-	_				aat Asn	_	_	_		_	-					153
					aaa Lys											201
		-			aac Asn 55											249
				-	ttg Leu	-				-	-		_			297
_					gag Glu	_							_	-	_	345
•		_	-		aaa Lys	_		_	555	_				_	-	393
	_			_	cca Pro		-						-		_	441
	-				tcg Ser 135	_		_		_						489
					ccg Pro											537
					aaa Lys											585
					cat His											633
					tat Tyr											681
					gac Asp 215											729

					ttt Phe									777
					gly ggg									825
					gct Ala									873
					ccg Pro									921
					gga Gly 295									969
					aat Asn									1017
					ggc Gly					_				1065
					acc Thr									1113
					act Thr									1161
				_	cgt Arg 375	_			_		_	-	_	1209
	ttt Phe	taag	gttt	ggt a	agggt	tttt	a gg	gcaag	ggctt	aga	agato	gaa		1254
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Ser Leu Asp Ile Ser Lys Lys Gly Ser Val Ile Leu Gly Ala Leu Leu
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Ser Ser Leu Trp Leu Thr Asn Pro Leu Asn Ala His Glu Lys Asn Gly
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Ala Phe Val Gly Ile Ser Leu Glu Val Gly Arg Ala Asp Gln Lys Thr
Asn Ala Tyr Lys Asn Gly Glu Leu Phe Gln Val Pro Phe Gly Asp Val
                                    90
Ser Ala Asn Asp Asp Gly Lys Val Pro Asp Gly Gln Thr Gly Gly Cys
                                105
Gln Pro Ala Ser Gly Thr Pro Gly Thr Pro Gly Tyr Thr Lys Ala Asn
                            120
                                                125
Cys Val Val Asn Trp Thr Ser Arg Thr Met Leu Ser Thr Asn Lys Asn
                        135
                                            140
Ile Pro Gly Arg Asn Gln Pro Met Tyr Gly Leu Gly Val Met Thr Gly
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                                        155
Tyr Lys His Phe Ile Gly Lys Lys Arg Trp Phe Gly Leu Arg Tyr Tyr
                                    170
                165
Gly Phe Phe Asp Tyr Gly His Thr Asn Phe Ser Asn Ser Arg Ala Ala
                               185
Asn Ala Ile Ser Pro Phe Tyr Leu Ser Asp Gln Lys Ala Asp Met Tyr
                            200
                                                205
Thr Tyr Gly Phe Gly Thr Asp Met Leu Phe Asn Ile Ile Asp Lys Pro
                        215
                                            220
Lys Ala Thr Ala Gly Phe Phe Leu Gly Val Asn Phe Ala Gly Asn Thr
                   230
                                       235
Trp Thr Asn Asn Arg Val Gly Tyr Phe Lys Asp Gly Tyr Val Tyr Gly
                                    250
                245
Val Asn Thr Asp Ala Asp Ala Tyr Met Thr Asn Ala Asp Gly Thr Ile
                                265
Thr Cys Gly Asp Thr Thr Pro Ala Ser Cys Asn Val Gly Ile Asn Pro
                            280
                                                285
Asn Ser Val Tyr Thr Thr Gly Lys Leu Asn Ala Lys Val Asn His Thr
                        295
Ile Phe Gln Phe Leu Val Asn Val Gly Ile Arg Thr Asn Ile Phe Glu
                    310
                                        315
His His Gly Ile Glu Phe Gly Ile Lys Ile Pro Thr Leu Pro Asn Tyr
                                    330
Phe Phe Lys Gly Ser Thr Thr Ile Arg Ala Lys Lys Gln Gly Pro Leu
           340
                                345
Glu Asn Gly Gln Pro Thr Thr Ile Thr Gly Ala Glu Thr Asn Phe Ser
                           360
Leu Thr Gln Thr Leu Arg Arg Gln Tyr Ser Met Tyr Leu Arg Tyr Val
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Tyr Thr Phe
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Ile Ile Phe Ile Gly Tyr Leu Phe Phe Asp Gly Val Ile Phe Pro Asn

25

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Val Leu Thr Gly Gly Met Met Leu Ser Gln Tyr Met Gly Gly Asp Lys
                    70
                                        75
Gly Trp Cys Glu Thr Pro Phe Gln Lys Ile Leu Met Leu Lys Val Ile
                                    90
                85
Leu Ala Leu Ser Ile Phe Leu Leu Val Leu Phe Ser Leu Ser Cys Lys
                                105
Phe Leu Gly Lys Lys Asn Pro Ile Gly Lys Tyr Ile His Pro Ile Ala
                            120
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Leu Thr Phe Gly Phe Leu Ile Ala Ile Leu Ala Lys Thr Met Trp Phe
                        135
Val
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                                                                   104
Leu Phe Glu Lys Ile Leu His Glu Met Arg Ser Arg Thr Ser Phe Leu
     5
                         10
ctt gct ttt gtc gtt tcg ctt att gtt ttt att ttt aat cta aaa ggg
                                                                   152
Leu Ala Phe Val Val Ser Leu Ile Val Phe Ile Phe Asn Leu Lys Gly
 20
                     25
gtt ttt caa ttg att ttt gag tct att ttc caa tac acc caa aac aaa
                                                                   200
Val Phe Gln Leu Ile Phe Glu Ser Ile Phe Gln Tyr Thr Gln Asn Lys
                 40
                                                          50
atc ctt tct ttt tct ctt tct ttt ttc ttt att ttc ttt ttc ttt tac
                                                                   248
Ile Leu Ser Phe Ser Leu Ser Phe Phe Phe Phe Phe Phe Tyr
             55
                                 60
gct att ttt ctt att ttt tat caa ata ttt ctt tgg tat ggg gct aaa
                                                                   296
Ala Ile Phe Leu Ile Phe Tyr Gln Ile Phe Leu Trp Tyr Gly Ala Lys
                             75
aaa tat aaa caa aat caa aga gat agt gaa att gtc tat aac att caa
                                                                   344
Lys Tyr Lys Gln Asn Gln Arg Asp Ser Glu Ile Val Tyr Asn Ile Gln
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										ctt Leu						440
										aga Arg						488
		_	_			-		_	_	gtc Val			-			536
										tat Tyr						584
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Met 1	Glu	Lys	Leu	Phe 5	Glu	Lys			10	Glu				15		
Met 1 Ser	Glu Phe	Lys Leu	Leu Leu 20	Phe 5 Ala	Glu Phe	Lys Val	Val	Ser 25	10 Leu	Ile	Val	Phe	Ile 30	15 Phe	Asn	
Met 1 Ser	Glu Phe	Lys Leu	Leu Leu 20	Phe 5 Ala	Glu Phe	Lys Val	Val	Ser 25	10 Leu		Val	Phe	Ile 30	15 Phe	Asn	
Met 1 Ser Leu	Glu Phe Lys	Lys Leu Gly 35	Leu Leu 20 Val	Phe 5 Ala Phe	Glu Phe Gln	Lys Val Leu	Val Ile 40	Ser 25 Phe	10 Leu Glu	Ile	Val Ile	Phe Phe 45	Ile 30 Gln	15 Phe Tyr	Asn Thr	
Met 1 Ser Leu Gln	Glu Phe Lys Asn 50	Lys Leu Gly 35 Lys	Leu Leu 20 Val Ile	Phe 5 Ala Phe Leu	Glu Phe Gln Ser	Lys Val Leu Phe 55	Val Ile 40 Ser	Ser 25 Phe Leu	10 Leu Glu Ser	Ile Ser	Val Ile Phe 60	Phe Phe 45 Phe	Ile 30 Gln Ile	15 Phe Tyr Phe	Asn Thr Phe	
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Met 1 Ser Leu Gln Phe 65 Gly	Glu Phe Lys Asn 50 Phe	Lys Leu Gly 35 Lys Tyr	Leu Leu 20 Val Ile Ala Lys	Phe 5 Ala Phe Leu Ile Tyr 85	Glu Phe Gln Ser Phe 70 Lys	Lys Val Leu Phe 55 Leu Gln	Val Ile 40 Ser Ile Asn	Ser 25 Phe Leu Phe Gln	10 Leu Glu Ser Tyr Arg 90	Ile Ser Phe Gln 75	Val Ile Phe 60 Ile Ser	Phe Phe 45 Phe Phe Glu	Ile 30 Gln Ile Leu Ile	15 Phe Tyr Phe Trp Val 95	Asn Thr Phe Tyr 80 Tyr	
Met 1 Ser Leu Gln Phe 65 Gly Asn	Glu Phe Lys Asn 50 Phe Ala Ile	Lys Leu Gly 35 Lys Tyr Lys Gln Lys	Leu Leu 20 Val Ile Ala Lys Lys 100	Phe 5 Ala Phe Leu Ile Tyr 85 Phe	Glu Phe Gln Ser Phe 70 Lys	Lys Val Leu Phe 55 Leu Gln Asn	Val Ile 40 Ser Ile Asn Glu Ile	Ser 25 Phe Leu Phe Gln Ile 105	10 Leu Glu Ser Tyr Arg 90 Lys	Ile Ser Phe Gln 75 Asp	Val Ile Phe 60 Ile Ser Glu	Phe Phe 45 Phe Phe Clu Leu Lys	Ile 30 Gln Ile Leu Ile Tyr 110	15 Phe Tyr Phe Trp Val 95 Arg	Asn Thr Phe Tyr 80 Tyr Cys	
Met 1 Ser Leu Gln Phe 65 Gly Asn Tyr	Glu Phe Lys Asn 50 Phe Ala Ile Ser Ile	Lys Leu Gly 35 Lys Tyr Lys Gln Lys 115	Leu Leu 20 Val Ile Ala Lys Lys 100 Lys	Phe 5 Ala Phe Leu Ile Tyr 85 Phe Gln	Glu Phe Gln Ser Phe 70 Lys Pro Asn	Lys Val Leu Phe 55 Leu Gln Asn Lys Leu	Val Ile 40 Ser Ile Asn Glu Ile 120	Ser 25 Phe Leu Phe Gln Ile 105 Leu	10 Leu Glu Ser Tyr Arg 90 Lys	Ile Ser Phe Gln 75 Asp	Val Ile Phe 60 Ile Ser Glu Lys Lys	Phe Phe 45 Phe Phe Glu Leu Lys 125	Ile 30 Gln Ile Leu Ile Tyr 110 Leu	15 Phe Tyr Phe Trp Val 95 Arg	Asn Thr Phe Tyr 80 Tyr Cys Asp	
Met 1 Ser Leu Gln Phe 65 Gly Asn Tyr Leu Phe	Glu Phe Lys Asn 50 Phe Ala Ile Ser Ile 130	Lys Leu Gly 35 Lys Tyr Lys Gln Lys 115 Asp	Leu Leu 20 Val Ile Ala Lys Lys 100 Lys	Phe 5 Ala Phe Leu Ile Tyr 85 Phe Gln Leu	Glu Phe Gln Ser Phe 70 Lys Pro Asn Asp	Lys Val Leu Phe 55 Leu Gln Asn Lys Leu 135	Val Ile 40 Ser Ile Asn Glu Ile 120 Ile	Ser 25 Phe Leu Phe Gln Ile 105 Leu Gly	10 Leu Glu Ser Tyr Arg 90 Lys Arg	Ile Ser Phe Gln 75 Asp Glu Thr Phe Lys	Val Ile Phe 60 Ile Ser Glu Lys Lys 140	Phe Phe 45 Phe Phe Glu Leu Lys 125 Ser	Ile 30 Gln Ile Leu Ile Tyr 110 Leu Arg	15 Phe Tyr Phe Trp Val 95 Arg Asp	Asn Thr Phe Tyr 80 Tyr Cys Asp Phe Arg	
Met 1 Ser Leu Gln Phe 65 Gly Asn Tyr Leu Phe 145	Glu Phe Lys Asn 50 Phe Ala Ile Ser Ile 130 Glu	Lys Leu Gly 35 Lys Tyr Lys Gln Lys 115 Asp	Leu Leu 20 Val Ile Ala Lys Lys 100 Lys Tyr	Phe 5 Ala Phe Leu Ile Tyr 85 Phe Gln Leu Lys	Glu Phe Gln Ser Phe 70 Lys Pro Asn Asp Asp	Lys Val Leu Phe 55 Leu Gln Asn Lys Leu 135 Asp	Val Ile 40 Ser Ile Asn Glu Ile 120 Ile Tyr	Ser 25 Phe Leu Phe Gln Ile 105 Leu Gly Ile	10 Leu Glu Ser Tyr Arg 90 Lys Arg Phe	Ile Ser Phe Gln 75 Asp Glu Thr	Val Ile Phe 60 Ile Ser Glu Lys Lys 140 Pro	Phe Phe 45 Phe Glu Leu Lys 125 Ser Asp	Ile 30 Gln Ile Leu Ile Tyr 110 Leu Arg Val	15 Phe Tyr Phe Trp Val 95 Arg Asp Asp	Asn Thr Phe Tyr 80 Tyr Cys Asp Phe Arg 160	

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Leu Gly Asp Phe Ala Arg Ile Gly Val Asn Ile His Ala Lys Lys Ala
         5
gtt tto aat aat gga ttt ttt aag att aaa aag ito tit ttg gat gtg
                                                                  152
Val Phe Asn Asn Gly Phe Phe Lys Ile Lys Lys Phe Phe Leu Asp Val
                         25
gtg gtg ggg gtc gtt ttc gct atc tgg ttt tat gta gtt aaa gat ttc
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Val Val Gly Val Val Phe Ala Ile Trp Phe Tyr Val Val Lys Asp Phe
                     40
acc cgt tgg cgt gtg gtg aat gat ttc atc gtt ttc taatttaaga
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Thr Arg Trp Arg Val Val Asn Asp Asp Phe Ile Val Phe
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Lys Ala Val Phe Asn Asn Gly Phe Phe Lys Ile Lys Lys Phe Phe Leu
Asp Val Val Gly Val Val Phe Ala Ile Trp Phe Tyr Val Val Lys
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Asp Phe Thr Arg Trp Arg Val Val Asn Asp Phe Ile Val Phe
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Glu Lys Ile Arg Lys Ile Leu Ala Asp Ile Glu Asp Ser Gln Asn Glu
att gaa atg ctt tta aaa tta gcg aat ttg agt ttg ggg gat ttt att
                                                                   152
Ile Glu Met Leu Leu Lys Leu Ala Asn Leu Ser Leu Gly Asp Phe Ile
gag att aaa aga ggg agc atg gac atg cca aag ggc gtg aat gaa gcg
                                                                   200
Glu Ile Lys Arg Gly Ser Met Asp Met Pro Lys Gly Val Asn Glu Ala
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Phe Phe Thr Gln Leu Ser Glu Glu Val Glu Arg Leu Lys Glu Leu Ile
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aac gct ttg aat aaa atc aaa aaa ggg tta ttg gtg ttt taaatgtgtg
                                                                   297
Asn Ala Leu Asn Lys Ile Lys Lys Gly Leu Leu Val Phe
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                                  75
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Phe Ile Glu Ile Lys Arg Gly Ser Met Asp Met Pro Lys Gly Val Asn
                            40
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ata ctc gct tgt aag gat ttt aaa gaa tcc gag ctc gct aaa gaa gtc 1 Ile Leu Ala Cys Lys Asp Phe Lys Glu Ser Glu Leu Ala Lys Glu Val 20 25 30	152
ata agc tat ttt aag cca aat acc aaa gcc att ctt ttc ccg gag ttt 2 Ile Ser Tyr Phe Lys Pro Asn Thr Lys Ala Ile Leu Phe Pro Glu Phe 35 40 45	200
agg gct aaa aaa aac gac gat ttg cgt tcg ttt ttt gaa gaa ttt tta 2 Arg Ala Lys Lys Asn Asp Asp Leu Arg Ser Phe Phe Glu Glu Phe Leu 50 55 60 65	248
cag ctt tta ggg ggt tta agg gag ttt tat caa gcc tta gaa aac aag 2 Gln Leu Leu Gly Gly Leu Arg Glu Phe Tyr Gln Ala Leu Glu Asn Lys 70 75 80	296
caa gaa act atc atc att gcc ccg att agc gcg tta ttg cac cct tta 3 Gln Glu Thr Ile Ile Ile Ala Pro Ile Ser Ala Leu Leu His Pro Leu 85 90 95	344
cct aaa aaa gaa ctt tta gaa agc ttt aaa atc act ctt tta gaa aaa 3 Pro Lys Lys Glu Leu Leu Glu Ser Phe Lys Ile Thr Leu Leu Glu Lys 100 105 110	392
tat aac ctt aag gat ttg aaa gac aag ctc ttt tat tat ggc tat gaa 4 Tyr Asn Leu Lys Asp Leu Lys Asp Lys Leu Phe Tyr Tyr Gly Tyr Glu 115 120 125	140
att tta gac tta gtg gaa gtg gaa ggc gaa gcg agc ttt agg ggg gat  Ile Leu Asp Leu Val Glu Val Glu Gly Glu Ala Ser Phe Arg Gly Asp  130 145	188
att gtg gat att tat gcg cca aat tct aaa gcg tat cgc ttg agt ttt 5 Ile Val Asp Ile Tyr Ala Pro Asn Ser Lys Ala Tyr Arg Leu Ser Phe 150 155 160	536
ttt gac acc gag tgt gag agc att aag gaa ttt gat ccc att act caa 5 Phe Asp Thr Glu Cys Glu Ser Ile Lys Glu Phe Asp Pro Ile Thr Gln 165 170 175	584
atg agc ctt aaa gaa gat ttg tta gaa att gaa atc ccc ccc acg ctt 6 Met Ser Leu Lys Glu Asp Leu Leu Glu Ile Glu Ile Pro Pro Thr Leu 180 185 190	532
ttt agt ttg gac gaa tca tct tat aag gat cta aaa aca aaa gtg gaa 6 Phe Ser Leu Asp Glu Ser Ser Tyr Lys Asp Leu Lys Thr Lys Val Glu	580

	195					200					205					
														ggt Gly		728
				-										aaa Lys 240		776
		-		-	-		_	-			_			agc Ser		824
														gaa Glu		872
				-	-							_		gāā Glu		920
		-	-							-			-	ccc Pro		968
														aac Asn 320		1016
														gat Asp		1064
													-	caa Gln		1112
														gtg Val		1160
					-	-					_	_		agc Ser	-	1208
												_		gaa Glu 400	-	1256
														tat Tyr		1304
gcg	caa	agc	gat	agc	gtg	cca	gct	aaa	gac	cgg	cta	ggg	aaa	agg	agc	1352

Ala	Gln	Ser 420	Asp	Ser	Val	Pro	Ala 425	Lys	Asp	Arg	Leu	Gly 430	Lys	Gly	Ser	
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-	_			-	tta Leu 455		-	-								1448
_	_	-			tta Leu											1496
		-			agc Ser	_		_	_	_		-	-			1544
					cac His											1592
					aca Thr	_			_					_		1640
					caa Gln 535											1688
					gag Glu											1736
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			_	_	tta Leu			_	_					_		1832
					aaa Lys											1880
					ggc Gly 615											1928
					ttt Phe											1976

		-		ctc Leu											2024
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-	-			aaa Lys	-				_	-		-	-		2120
				atc Ile 695					_	_				_	2168
	-		_	gat Asp								-		_	2216
	_			gct Ala		-	_	-	_		_		-		2264
				cag Gln											2312
		_		aac Asn	_		_				-				2360
				gat Asp 775											2408
		_		ttt Phe	_					_	_			_	2456
				tta Leu											2504
				gag Glu											2552
				ctc Leu											2600
				tat Tyr 855											2648

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ggc gtg agc gc Gly Val Ser Al 88	a Phe Leu		Leu Ile Ala		
aga ttg gat tt Arg Leu Asp Le 900					
gtg ggg caa at Val Gly Gln Il 915	e His Glu				
gat ttg agc gc Asp Leu Ser Al 930					
aac cag ctt gg Asn Gln Leu Gl					
atc act tat ag Ile Thr Tyr Se 96	r Asp Glu	-	Ser Leu Lys	•	
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Leu Gln Leu Le 65	Gly Gly 70	Leu Arg Glu	Phe Tyr Gln 75		Asn 80

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Leu Pro Lys Lys Glu Leu Leu Glu Ser Phe Lys Ile Thr Leu Leu Glu
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Lys Tyr Asn Leu Lys Asp Leu Lys Asp Lys Leu Phe Tyr Tyr Gly Tyr
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Glu Ile Leu Asp Leu Val Glu Val Glu Gly Glu Ala Ser Phe Arg Gly
                        135
Asp Ile Val Asp Ile Tyr Ala Pro Asn Ser Lys Ala Tyr Arg Leu Ser
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Phe Phe Asp Thr Glu Cys Glu Ser Ile Lys Glu Phe Asp Pro Ile Thr
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Gln Met Ser Leu Lys Glu Asp Leu Leu Glu Ile Glu Ile Pro Pro Thr
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Leu Phe Ser Leu Asp Glu Ser Ser Tyr Lys Asp Leu Lys Thr Lys Val
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Glu Gln Ser Pro Leu Asn Ser Phe Ser Lys Asp Leu Thr Ser Phe Gly
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Leu Trp Phe Leu Gly Glu Lys Ala Gln Asp Leu Leu Ile Val Tyr Lys
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Ser Ile Ile Ser Pro Arg Ala Leu Glu Glu Ile Gln Glu Leu Ala Ser
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Leu Asn Glu Leu Asp Cys Glu Arg Phe Lys Phe Leu Lys Val Leu Glu
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Asn Ala Gln Gly Tyr Glu Asp Leu Glu Ile His Ala His Ala Leu Glu
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Gly Phe Ile Ala Leu His Ser Asn His Lys Ile Thr Leu Leu Ala Pro
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Asn Lys Thr Ile Leu Asp Asn Ala Ile Ser Ala Leu Asp Ala Gly Asn
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Met Glu Cys Val Ile Ala Pro Phe Val Leu Asn Phe Lys Thr Pro Asp
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Gly Ile Phe Ile Ser Leu Asn Ser Phe Glu Arg Lys Lys Arg Gln
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Lys Ser Lys Leu Ala Leu Asn Glu Leu Asn Pro Gly Glu Trp Val Val
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His Asp Asp Tyr Gly Val Gly Val Phe Ser Gln Leu Val Gln His Ser
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                                            380
Val Leu Gly Ser Lys Arg Asp Phe Leu Glu Ile Ala Tyr Leu Gly Glu
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Asp Lys Leu Leu Pro Val Glu Asn Leu His Leu Ile Ala Arg Tyr
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Val Ala Gln Ser Asp Ser Val Pro Ala Lys Asp Arg Leu Gly Lys Gly
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Ser Phe Leu Lys Leu Lys Ala Lys Val Arg Thr Lys Leu Leu Glu Ile
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Ala Ser Lys Ile Ile Glu Leu Ala Ala Glu Arg Asn Leu Ile Leu Gly
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Lys Lys Met Asp Val His Leu Ala Glu Leu Glu Val Phe Lys Ser His
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Asp Val Gly Phe Gly Lys Thr Glu Val Ala Met His Ala Ile Phe Cys
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Arg Thr Leu Asn Met Ala Leu Ser Gln Ile Lys Gly Ile Ser Ser Leu
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Lys Thr Pro Pro Thr Asp Arg Lys Pro Ser Arg Thr Phe Leu Lys Glu
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Lys Asn Asp Glu Leu Lys Glu Ile Ile Tyr Arg Glu Leu Arg Arg
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Val Lys Thr Lys Leu Glu Asp Leu Ile Pro Lys Leu Lys Ile Ala Ile
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Leu His Ser Gln Ile Asn Ala Asn Glu Ser Glu Glu Ile Met Leu Glu
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Phe Ala Lys Gly Asn Tyr Gln Val Leu Leu Cys Thr Ser Ile Val Glu
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Ser Gly Ile His Leu Pro Asn Ala Asn Thr Ile Ile Ile Asp Asn Ala
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Gln Asn Phe Gly Leu Ala Asp Leu His Gln Leu Arg Gly Arg Val Gly
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Arg Gly Lys Lys Glu Gly Phe Cys Tyr Phe Leu Ile Glu Asp Gln Lys
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Ser Leu Asn Glu Gln Ala Leu Lys Arg Leu Leu Ala Leu Glu Lys Asn
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Ser Tyr Leu Gly Ser Gly Glu Ser Val Ala Tyr His Asp Leu Glu Ile
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Arg Gly Gly Asn Leu Leu Gly Gln Asp Gln Ser Gly His Ile Lys
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Asn Ile Gly Tyr Ala Leu Tyr Thr Arg Met Leu Glu Asp Ala Ile Tyr
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Glu Leu Ser Gly Gly Lys Lys Arg Leu Glu Lys Ser Val Glu Ile Gln
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Asp Asp Leu Ser Ala Gln Phe Leu Gln Ile Ile Thr Leu Lys Ile Leu
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Thr Ile Thr Tyr Ser Asp Glu Lys Lys Glu Ser Leu Lys Ala Pro Ser

gac aca age gaa tet ttt cat ttg agt ggt gtg ata agt tta gta tea

Asp	Thr	Ser 135	Glu	Ser	Phe	His	Leu 140	Ser	Gly	Val	Ile	Ser 145	Leu	Val	Ser	
					GJA aaa											537
					caa Gln 170											585
_	_				aaa Lys	_										633
			-		ccc Pro				-	-						681
	-	_		_	gat Asp	_	_			_						729
	_				caa Gln	_				_				_	_	777
					atc Ile 250											825
_		_			gaa Glu		_		~	•						873
					ttc Phe		-						_			921
					act Thr											969
					gaa Glu											1017
					cgt Arg 330			_		-		_			_	1065
					tac Tyr											1113

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          Met Gly Asn Val Asn Pro Pro Lys Thr Ser Tyr Lys Glu Ser
ctt tct acc att gcg agc att gaa tac gct atc gcg cat gtg ggc gtt
                                                                   339
Leu Ser Thr Ile Ala Ser Ile Glu Tyr Ala Ile Ala His Val Gly Val
 15
                     20
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Gln Asn Leu Ile Ile Cys Gly His Ser Asp Cys Gly Ala Cys Gly Ser
                 35
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gtt cat tta atc cat gat gaa acc acc aaa gct aaa acc cct tac att
                                                                   435
Val His Leu Ile His Asp Glu Thr Thr Lys Ala Lys Thr Pro Tyr Ile
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gca aac tgg ata caa ttt tta gag cct gtt aaa gaa gag tta aaa aac
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Ala Asn Trp Ile Gln Phe Leu Glu Pro Val Lys Glu Glu Leu Lys Asn
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cac ccg caa ttc agc aac cat ttc gcc aag cgt tca tgg ctt aca gag
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His Pro Gln Phe Ser Asn His Phe Ala Lys Arg Ser Trp Leu Thr Glu
cgt ttg aat gcg cgc ttg caa ctc aac aac ctc tta agc tat gat ttc
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Arg Leu Asn Ala Arg Leu Gln Leu Asn Asn Leu Leu Ser Tyr Asp Phe
 95
                    100
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Ile Gln Glu Lys Ala Ser Lys Asn Glu Leu Lys Ile Phe Gly Trp His
                115
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tac atc ata gaa aca ggc agg att tat aat tat aat ttt gaa agc cat
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Tyr Ile Ile Glu Thr Gly Arg Ile Tyr Asn Tyr Asn Phe Glu Ser His

130 135 140 723 ttt ttt gag ccg att gga gaa acc att aaa caa agg aaa agt cat gaa Phe Phe Glu Pro Ile Gly Glu Thr Ile Lys Gln Arg Lys Ser His Glu 150 155 aac ttc taaaacaaaa acccctaaat ccgttttaat cgctgggcca tgcgtcattg 779 Asn Phe 160 780 а <210> 230 <211> 160 <212> PRT <213> Helicobacter pylori <400> 230 Met Gly Asn Val Asn Pro Pro Lys Thr Ser Tyr Lys Glu Ser Leu Ser Thr Ile Ala Ser Ile Glu Tyr Ala Ile Ala His Val Gly Val Gln Asn 25 . Leu Ile Ile Cys Gly His Ser Asp Cys Gly Ala Cys Gly Ser Val His 40 Leu Ile His Asp Glu Thr Thr Lys Ala Lys Thr Pro Tyr Ile Ala Asn 55 Trp Ile Gln Phe Leu Glu Pro Val Lys Glu Glu Leu Lys Asn His Pro Gln Phe Ser Asn His Phe Ala Lys Arg Ser Trp Leu Thr Glu Arg Leu 90 Asn Ala Arg Leu Gln Leu Asn Asn Leu Leu Ser Tyr Asp Phe Ile Gln 105 110 Glu Lys Ala Ser Lys Asn Glu Leu Lys Ile Phe Gly Trp His Tyr Ile 120 Ile Glu Thr Gly Arg Ile Tyr Asn Tyr Asn Phe Glu Ser His Phe Phe 135 140 Glu Pro Ile Gly Glu Thr Ile Lys Gln Arg Lys Ser His Glu Asn Phe 145 150 155

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cat ttt tta aaa atc gtt ttt tta gta ggc atg tgc gtt tca agt ttg
                                                                   163
His Phe Leu Lys Ile Val Phe Leu Val Gly Met Cys Val Ser Ser Leu
                         10
                                                                   211
ttc gct gaa ggt tta gag ggg ttt ttt aac gcc cta gaa gcc cag ctc
Phe Ala Glu Gly Leu Glu Gly Phe Phe Asn Ala Leu Glu Ala Gln Leu
                                         30
                     25
                                                                   259
aaa agc ccc atc gct aag ggg att tta atg gtg att ttc ata ggg atc
Lys Ser Pro Ile Ala Lys Gly Ile Leu Met Val Ile Phe Ile Gly Ile
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gct att tat gtg tgg agg aat ttg gac cgg tgg aaa gag atc tta ttc
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Ala Ile Tyr Val Trp Arg Asn Leu Asp Arg Trp Lys Glu Ile Leu Phe
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acg atc ctt ggc gtg gtg ttt ggg att ttt tta ttc ttt aaa gct ccg
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Thr Ile Leu Gly Val Val Phe Gly Ile Phe Leu Phe Phe Lys Ala Pro
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                             75
agt tta gcg aat tgg ttt atg gga att ttt taatgattat cctgtcagcg
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Ser Leu Ala Asn Trp Phe Met Gly Ile Phe
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Ile Gly Ile Ala Ile Tyr Val Trp Arg Asn Leu Asp Arg Trp Lys Glu
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_						att Ile										400
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						ttg Leu										496
						agc Ser										544
						tta Leu										592
Gly	gtg Val	aaa Lys 175	gag Glu	cat His	ggc Gly	tta Leu	gag Glu 180	Gly	gtg Val	agc Ser	gly	ttg Leu 185	cat His	ttc Phe	cat His	640
						gct Ala 195										688
gta	gaa	aag	cat	ttc	agg	ccc	tat	tta	gaa	aac	atg	gcg	tgg	gtg	aat	736

Val 205	Glu	Lys	His	Phe	Arg 210	Pro	Tyr	Leu	Glu	Asn 215	Met	Ala	Trp	Val	Asn 220	
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					aaa Lys 370											1216
					gat Asp											1264
					aaa Lys				taaa	agctt	tt g	gatta	aggo	et		1311
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tgattaacgc ctatgacaac accatttttt acaacgacta tctgctagat cgaaagatca 240
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      Met Leu Glu Asn Ala Lys Gln Pro Ala Leu Met Ile Tyr Leu Ser
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gat cat ggc gaa agt ttg ggc gaa gaa gcg ttc tat ttg cat ggc att
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Asp His Gly Glu Ser Leu Gly Glu Glu Ala Phe Tyr Leu His Gly Ile
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cct aaa agc atc gcc ccc aaa gaa caa tac gag atc ccc ttt atc gtt
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Pro Lys Ser Ile Ala Pro Lys Glu Gln Tyr Glu Ile Pro Phe Ile Val
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                                  40
                                                      45
tat gct aat gag cct ttc aaa gaa aag cat tcc atc att caa acc caa
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Tyr Ala Asn Glu Pro Phe Lys Glu Lys His Ser Ile Ile Gln Thr Gln
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                                                  60
acc ccc att aat caa aat gtg att ttc cat agc gtt tta ggg gtg ttt
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Thr Pro Ile Asn Gln Asn Val Ile Phe His Ser Val Leu Gly Val Phe
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                                         90
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-	_			-						acg Thr 170			-	_	-	647
		-								aac Asn			_			695
										ggc Gly						743
										gaa Glu						791
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Gln	Gly 50	Ile	Asp	Phe	Gly	Glu 55	Trp	Asp	Lys	Asn	Phe 60	Lys	Leu	Leu	Glu	
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Ile	Phe	Cys	Ser	Tyr 85	Arg	Phe	Ile	Ser	Tyr 90	Ile	Ala	Asp	Phe	Leu 95		
Glu	Asn	Gly	Phe 100		Val	Lys	Asp	Phe 105		Gln	Trp	Val	Lys 110		Asn	
Pro	Met	Pro 115		Asn	Ile	His	Arg 120		Tyr	Val	Gln	Asp 125		Glu	Phe	

Ala Leu Trp Ala Val Lys Lys Ala Lys Trp Val Phe Asn Lys Pro 130 135 Lys Asn Glu Lys Tyr Leu Arg Pro Leu Ile Leu Lys Ser Pro Val Val 160 150 155 Ser Gly Leu Glu Lys Thr Lys His Pro Thr Gln Lys Ser Leu Ala Leu 170 Met Glu Lys Ile Ile Ser Ile His Thr Asn Pro Asn Asp Ile Val Leu 185 190 Asp Pro Phe Met Gly Ser Gly Thr Thr Gly Leu Ala Cys Lys Asn Leu 205 200 Glu Arg Asn Phe Ile Gly Ile Glu Ser Glu Lys Glu Tyr Phe Gln Thr 215 220 Ala Lys Lys Arg Leu Asn Leu Phe 230 <210> 239 <211> 1181 <212> DNA <213> Helicobacter pylori <220> <221> CDS <222> (60)...(1124) <400> 239 ttgcgttttg aaaccgatga tttttcaacg cttattgatc gtatttgtga aagcttgaa atg aat tat aaa att tta gat tta ttt tgt ggg gct ggg ggt ttt agc 107 Met Asn Tyr Lys Ile Leu Asp Leu Phe Cys Gly Ala Gly Gly Phe Ser 15 gct ggg tta gag tgt tta gaa gag ttt gac gct tta ata ggg cta gat 155 Ala Gly Leu Glu Cys Leu Glu Glu Phe Asp Ala Leu Ile Gly Leu Asp 20 25 tgc gat aaa caa gcc cta atc act ttt gaa aac aac cat aaa aac gcc 203 Cys Asp Lys Gln Ala Leu Ile Thr Phe Glu Asn Asn His Lys Asn Ala 40 ata ggc gtt tgt ggg gac atc act caa acc gaa att aaa gaa aaa gtc 251 Ile Gly Val Cys Gly Asp Ile Thr Gln Thr Glu Ile Lys Glu Lys Val 50 55 atc aaa cta gct aaa aaa tta gaa atc aac atg atc att ggc ggg cct 299 Ile Lys Leu Ala Lys Lys Leu Glu Ile Asn Met Ile Ile Gly Gly Pro 65 70 75 80 cca tgt caa ggc ttt tct aat aaa ggg aaa aat tta ggg cta aaa gac 347 Pro Cys Gln Gly Phe Ser Asn Lys Gly Lys Asn Leu Gly Leu Lys Asp 85 90 95 cct agg aat ttt tta ttc tta gaa tat ata gaa ata gtc aaa gcc ata 395 Pro Arg Asn Phe Leu Phe Leu Glu Tyr Ile Glu Ile Val Lys Ala Ile 100 105 110

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											gat Asp				539
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											aac Asn				779
_	_										caa Gln				827
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Ile Lys Leu Ala Lys Lys Leu Glu Ile Asn Met Ile Ile Gly Gly Pro
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Pro Cys Gln Gly Phe Ser Asn Lys Gly Lys Asn Leu Gly Leu Lys Asp
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Ala Lys Gly Tyr Phe Leu Glu Glu Ile Lys Glu Arg Leu Asn Ala Leu
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Pro Gln Asn Arg Glu Arg Ala Phe Ile Val Gly Ala Ser Arg Phe Ser
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Phe Asp Phe Asn Leu Leu Glu Pro Ser Gln Ser Val Asn Val Gln Asp
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Lys Asp Ser Pro Lys Leu Tyr Asn His Gln Ala Thr Asn His Ser Gln
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Cys Leu Pro Lys Asn Leu His Gly Lys Gln Gln Phe Lys Ser Thr Trp
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Gly Arg Leu Asn Trp Asn Lys Ile Ser Pro Thr Ile Asp Thr Arg Phe
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Asp Thr Pro Ser Asn Gly Thr Asn Ser His Pro Glu Leu His Arg Ser
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					ttt Phe							537
					aaa Lys 170							585
					gct Ala							633
				_	ttt Phe							681
	_				ata Ile							729
_			_	-	ata Ile		_					777
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			_	_	aaa Lys	_		_	 _		_	921
					ccc Pro		_				•	969
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Gln Ala Thr Gln Ile His His Ile Phe Pro Ile Gln Asp Phe Pro Ile
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Ile Ala Asn Tyr Ile Glu Asn Leu Ile Ala Leu Thr Pro Asn Gln His
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Phe Ile Tyr Ala His Pro Asn Asn Gln Thr Arg Leu Ile Asp Lys Asp
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Leu Lys Val Ile Asp Ala Phe Tyr Phe Asp Phe Asn Lys Ser Lys Asp
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Tyr Ile Ser Leu Phe Ser Gly Ala Gly Val Gly Cys Tyr Gly Leu Leu
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Leu Asn Ile Gln Arg Ile Asn Arg Lys Cys Lys Leu Asp Glu Ser Tyr
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				cca Pro												401
_			_	gag Glu						_		_	_	_		449
_	-			caa Gln 125				_					_		_	497
	_			aaa Lys			_		_			-		_		545
-				atg Met					_	_		_		_		593
				atc Ile												641
				gtg Val												689
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				tcg Ser												785
				cat His	-		_				-		_		_	833
				tta Leu												881
				cct Pro												929

-											aaa Lys					977
_		_				-		-		_	gct Ala	_			_	1025
											aga Arg					1073
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	-				_	_	_				gaa Glu 420				_	1361
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				_							ttt Phe				~	1457
				_		_					agt Ser	_				1505
_					_					-	gct Ala					1553
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		gtg Val	-						_					_		1745
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		gaa Glu	_					_					_	_	_	1841
		aac Asn									_		_		_	1889
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		gtt Val 635														2033
		agc Ser														2081
		gct Ala														2129
		gtg Val														2177
		att Ile														2225
		agc Ser														2273

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155

Arg Phe Phe Ile Leu Glu Asn Val Pro Ser Phe Tyr Lys Thr Gly Cys

Ile Asp Lys Asn Asp Asn Leu Leu Glu Ile Gly Ser Met Ile Glu Gln

Asn Leu Ser Gly Asp Tyr Met Leu Tyr Asp Glu Val Ile Asn Phe Lys

135

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Phe	Lys 210	Gln	Glu	Lys	Thr	Leu 215	Lys	Glu	Val	Ile	Gly 220	Ser	Leu	Lys	Pro
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Thr	Tyr	Pro	Lys	His 245	Met	Gln	Glu	Trp	Ile 250	Lys	Asp	Leu	Lys	Glu 255	Gly
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	530				Tyr	535					540				
545					Lys 550					555					560
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Gln Lys Ile Ile Asp Asp Ser Leu Glu Leu Ala Lys Lys Leu Gln Asp
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Ser Ile Ser Asn His Leu Ser Asp Gln Glu Lys Ala Phe His Ser Lys
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                             25
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													att Ile			296
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													aga Arg			392
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			_			_				_		-	aat Asn			488
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			_										tcc Ser 175			584
		-									_		gaa Glu			632
				_		_		_	_			_	ctg Leu	_		680
							-						ttg Leu	_	_	728
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													tct Ser			872

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	-		_					_		_	tat Tyr	_		_	_	1112
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475

1496

1544

Thr Leu Asp Asn Ala Thr His Arg Thr Gln Asp Arg Asn Ala Lys Gln

agc ggg cat acc act tac cca aac cac tcc ttt aaa aac gaa agc gat

Ser Gly His Thr Thr Tyr Pro Asn His Ser Phe Lys Asn Glu Ser Asp

acc gat ttt att tta aaa gcc aac cga gaa tgg gct aaa aaa gtg cgc

455

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			-	_	cct Pro						-			-	-	1640
					atc Ile 535	_	-					_	_	_		1688
					gaa Glu											1736
					gaa Glu			_		_	-			_		1784
					aga Arg		_	_					-		_	1832
					gct Ala											1880
					tac Tyr 615											1928
					ttc Phe							-	-		_	1976
					gtg Val											2024
					cgg Arg											2072
					ctt Leu											2120
					tac Tyr 695											2168

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						cct Pro						_	2504
						gaa Glu							2552
						gaa Glu 840							2600
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						ctt Leu							2696
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_		-	-		Leu					Lys	gtt Val		-		Gly	3176
				Glu	_	-			Ser	_	atg Met			Val	-	3224
	_	_	Leu					Pro		_	ctc Leu	_	Ile	_		3272
_	_	Thr			gat Asp						gaa	_				3320
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	Gln	gca				1080 att Ile	gtt	tat	gaa	agc	1085 ttg Leu	caa	aaa	ttt	agc	3368
Leu 1090 gag	Gln ) aaa	gca Ala ttg	Ser	Ala	Pro 1095 ttc Phe	att Ile aat	gtt Val	tat Tyr gtc	gaa Glu cgt	agc Ser 1100 tat Tyr	1085 ttg Leu	caa Gln	aaa Lys agc	ttt Phe	agc Ser 1105 ttg Leu	3368 3416
Leu 1090 gag Glu gat	Gln aaa Lys tta	gca Ala ttg Leu	Ser aat Asn	Ala act Thr 1110 gaa Glu	Pro 1095 ttc Phe caa	att Ile aat Asn	gtt Val cgt Arg	tat Tyr gtc Val	gaa Glu cgt Arg 1115 tta Leu	agc Ser 1100 tat Tyr	ttg Leu ctc	caa Gln aaa Lys	aaa Lys agc Ser	ttt Phe gat Asp 1120 gcc Ala	agc Ser 1105 ttg Leu	

1140 1145 1150

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Asp	Thr	Lys 115		Met	Val	Leu	Asp		Glu	Glu	Ser	Gln 125		Lys	Glu	
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	Tyr	Ser	Gln	Ala 165		Lys	Ser	Asn	Tyr 170		Gln	Tyr	Ile	Ser		
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Gln Leu Phe Arg Glu Arg Arg Gly Asp Leu Ile Gly Ile Ser Ala Leu
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aag aad Lys Asi 5															163
aaa cgo Lys Aro 20							_				_				211
cct tta Pro Le											_	_		-	259
gaa tti Glu Phe															307
gca cad Ala His															355
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693
Lys Lys Pro Asp Phe Asp Glu Glu Asn Phe
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Lys Pro Ser Ala His Glu Ile Phe Ser Leu Cys Lys Glu Leu Lys Asn
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Gln Ser Gln Glu Lys Tyr Lys Arg Glu Arg Gly Glu Thr Leu Glu
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					gaa Glu											396
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					caa Gln 135											492
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					gat Asp											588
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tct ggt gtg cga gaa tgc att gaa gaa agc gaa ggg gtg gat ggg gc Ser Gly Val Arg Glu Cys Ile Glu Glu Ser Glu Gly Val Asp Gly Ala 210 215 220	ı
gtg agt gaa acc gct agt tct cat tta tgc gtg aaa gct tta gcg aaa Val Ser Glu Thr Ala Ser Ser His Leu Cys Val Lys Ala Leu Ala Lys 230 235 240	
ggc tca gaa ccc tta ttg cat tta aga gaa aaa atc gct cgc ttg gt Gly Ser Glu Pro Leu Leu His Leu Arg Glu Lys Ile Ala Arg Leu Va 245 250 255	
acg caa acc acc acg caa aag gtt tgaaagcact tcaaaaagat taaagtcc Thr Gln Thr Thr Gln Lys Val 260 265	t 882
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Gly Ala Asp Gly Arg Cys Val Ile Glu Asp Asn Phe Phe Thr Pro Pro 20 25 30  Phe Lys Leu Met Ala Pro Phe Tyr Pro Lys Asp Asp Leu Ala Glu Ile	
Gly Ala Asp Gly Arg Cys Val Ile Glu Asp Asn Phe Phe Thr Pro Pro 20 25 30  Phe Lys Leu Met Ala Pro Phe Tyr Pro Lys Asp Asp Leu Ala Glu Ile 35 40 45  Met Leu Leu Ala Val Ser Pro Gly Met Met Arg Gly Asp Ala Gln Asp	2
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Gly Ala Asp Gly Arg Cys Val Ile Glu Asp Asn Phe Phe Thr Pro Pro 20 25 30  Phe Lys Leu Met Ala Pro Phe Tyr Pro Lys Asp Asp Leu Ala Glu Ile 35 40 45  Met Leu Leu Ala Val Ser Pro Gly Met Met Arg Gly Asp Ala Gln Asp 50 55 60  Val Gln Leu Asn Ile Gly Pro Asn Cys Lys Leu Arg Ile Thr Ser Glr 65 70 75 80  Ser Phe Glu Lys Ile His Asn Thr Glu Asp Gly Phe Ala Ser Arg Asp	
Gly Ala Asp Gly Arg Cys Val Ile Glu Asp Asn Phe Phe Thr Pro Pro 20 25 30  Phe Lys Leu Met Ala Pro Phe Tyr Pro Lys Asp Asp Leu Ala Glu Ile 35 40 45  Met Leu Leu Ala Val Ser Pro Gly Met Met Arg Gly Asp Ala Gln Asp 50 55 60  Val Gln Leu Asn Ile Gly Pro Asn Cys Lys Leu Arg Ile Thr Ser Glr 65 70 75 80  Ser Phe Glu Lys Ile His Asn Thr Glu Asp Gly Phe Ala Ser Arg Asp 85 90 95  Met His Ile Val Val Gly Glu Asn Ala Phe Leu Asp Phe Ala Pro Phe	
Gly Ala Asp Gly Arg Cys Val Ile Glu Asp Asn Phe Phe Thr Pro Pro 20 25 30  Phe Lys Leu Met Ala Pro Phe Tyr Pro Lys Asp Asp Leu Ala Glu Ile 35 45  Met Leu Leu Ala Val Ser Pro Gly Met Met Arg Gly Asp Ala Gln Asp 50 55 60  Val Gln Leu Asn Ile Gly Pro Asn Cys Lys Leu Arg Ile Thr Ser Glr 65 70 75 80  Ser Phe Glu Lys Ile His Asn Thr Glu Asp Gly Phe Ala Ser Arg Asp 85 90 95  Met His Ile Val Val Gly Glu Asn Ala Phe Leu Asp Phe Ala Pro Phe 100 100 105 110  Pro Leu Ile Pro Phe Glu Asn Ala His Phe Lys Gly Asn Thr Thr Ile	
Gly Ala Asp Gly Arg Cys Val Ile Glu Asp Asn Phe Phe Thr Pro Pro 20	
Gly Ala Asp Gly Arg Cys Val Ile Glu Asp Asn Phe Phe Thr Pro Pro 20	
Gly Ala Asp Gly Arg Cys Val Ile Glu Asp Asn Phe Phe Thr Pro Pro 20	
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210 Ala Val Ser	Glu Thr Ala	215 Ser Ser H:	is Leu Cve	220 Val Lvs Ala	Leu Ala
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	Glu Pro Leu 245		250	Lys lie Ala	255
Val Thr Gln	Thr Thr Thr 260	_	al 65		
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	aga aaa gaa Arg Lys Glu	_	-		
	aga ttg ggc Arg Leu Gly				
	att tat ggc Ile Tyr Gly 40				
	ggc atg agc Gly Met Ser 55				
	atc act aac Ile Thr Asn 70	Ala Leu Il			
	att ggt att Ile Gly Ile				
	aaa gac atg Lys Asp Met				
	act gaa gcc Thr Glu Ala 120			-	_

ggt Gly											488
gct Ala											536
gct Ala											584
aaa Lys 180	 _	-									632
ttg Leu											680
gaa Glu											728
cct Pro	_				_	_	-	_		_	776
caa Gln											824
gac Asp 260		-	_		-		-				872
gaa Glu	 -	 		_		•			_	_	920
 gaa Glu			-								968
gtg Val											1016
ttg Leu											1064
cgc Arg 340											1112

					tct Ser										1160
					tgg Trp										1208
					gaa Glu										1256
					tac Tyr										1304
	_		_		tct Ser 425	_	_					_	_	-	1352
_	_	 -		_	ttc Phe		-	_				_			1400
					ttg Leu	_		_		_			_		1448
					gtt Val			_	_	_		_			1496
					gca Ala										1544
					gaa Glu 505	_				_	_			-	1592
					aac Asn										1640
		_			gaa Glu	_			_						1688
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Lys Thr Leu Arg Glu Gly Met Ser Gln Ser Asn Asn Pro Ser Lys Glu
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Glu Leu Asp Leu Ile Ile Thr Asn Ala Leu Ile Val Asp Tyr Thr Gly
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Ser Val Gly Pro Ala Thr Glu Ala Leu Ala Gly Glu Gly Leu Ile Val
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Thr Gly Pro Ala Asp Gly Thr Asn Ala Thr Thr Ile Thr Pro Gly Arg
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Arg Asn Leu Lys Trp Met Leu Arg Ala Ala Glu Glu Tyr Ser Met Asn
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Asp Gln Ile Glu Ala Gly Ala Ile Gly Phe Lys Ile His Glu Asp Trp
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Gly Thr Thr Pro Ser Ala Ile Asn His Ala Leu Asp Val Ala Asp Lys
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Tyr Asp Val Gln Val Ala Ile His Thr Asp Thr Leu Asn Glu Ala Gly
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Cys Val Glu Asp Thr Met Ala Ala Ile Ala Gly Arg Thr Met His Thr
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Phe His Thr Glu Gly Ala Gly Gly Gly His Ala Pro Asp Ile Ile Lys
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Val Ala Gly Glu His Asn Ile Leu Pro Ala Ser Thr Asn Pro Thr Ile
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                                            300
Pro Phe Thr Val Asn Thr Glu Ala Glu His Met Asp Met Leu Met Val
                    310
                                        315
Cys His His Leu Asp Lys Ser Ile Lys Glu Asp Val Gln Phe Ala Asp
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Ser Arg Ile Arg Pro Gln Thr Ile Ala Ala Glu Asp Thr Leu His Asp
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Met Gly Ile Phe Ser Ile Thr Ser Ser Asp Ser Gln Ala Met Gly Arg
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Lys Glu Phe Gly Arg Leu Lys Glu Glu Lys Gly Asp Asn Asp Asn Phe
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Arg Ile Lys Arg Tyr Leu Ser Lys Tyr Thr Ile Asn Pro Ala Ile Ala
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His Gly Ile Ser Glu Tyr Val Gly Ser Val Glu Val Gly Lys Val Ala
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                                                     430
Asp Leu Val Leu Trp Ser Pro Ala Phe Phe Gly Val Lys Pro Asn Met
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Ile Ile Lys Gly Gly Phe Ile Ala Leu Ser Gln Met Gly Asp Ala Asn
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Ala Ser Ile Pro Thr Pro Gln Pro Val Tyr Tyr Arg Glu Met Phe Ala
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                                         475
His His Gly Lys Ala Lys Tyr Asp Ala Asn Ile Thr Phe Val Ser Gln
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Ala Ala Tyr Asp Lys Gly Ile Lys Glu Glu Leu Gly Leu Glu Arg Gln
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                                505
Val Leu Pro Val Lys Asn Cys Arg Asn Ile Thr Lys Lys Asp Met Gln
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Phe Asn Asp Thr Thr Ala His Ile Glu Val Asn Pro Glu Thr Tyr His
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                                                                   104
Ile Ala His Ser Asn Glu Ile Ala Arg Pro Ile Phe Lys Ser Gln Asp
cag ctt ttc act ctt tat caa ggg gat tgt aat gag gtt ttg ccc caa
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Gln Leu Phe Thr Leu Tyr Gln Gly Asp Cys Asn Glu Val Leu Pro Gln
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ttt gaa aac cag ttt gat ttg att ttt gct gat ccg cct tat ttc ctc
                                                                   200
Phe Glu Asn Gln Phe Asp Leu Ile Phe Ala Asp Pro Pro Tyr Phe Leu
 35
                     40
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360

355

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											gag Glu		296
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_		-								_	gly	-	392
_		_	_								tgg Trp		440
					_	_	_		_	_	cat His 145		488
				 _		_	_			_	cat His	_	536
		 _		_				_	_		atg Met	_	584
											aat Asn		632
											ctt Leu		680
											ggg Gly 225		728
											att Ile		776
											aaa Lys		824
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			gaa Glu 15													339
			gac Asp													387
			ctc Leu													435
			tgg Trp													483
			ccc Pro													531
			gtg Val 95								-				_	579
	aac Asn		taaa	agtg	gct t	caaaa	aaaaq	gg ct	ttat	tggt	ggt :	ttaa	aatc			628
tttg	gattt	ta q	gatco	ggat												646
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	)> 25															
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	Glu	Phe	Thr 20	_	Asn	Leu	Asp	Leu 25		Tyr	Pro	Phe	Met 30		Met	
Thr	Thr	Arg 35	Asp	Lys	Glu	Glu	Glu 40		Tyr	Trp	Asp	Met 45		Leu	Met	

Ala Ala Tyr Leu Ile Glu Asn Pro Glu Lys Ile Ile Gly Pro Lys 75 70 His Trp Leu Phe Gly His Glu Asn Ile Leu Cys Lys Glu Trp Val Lys Ile Glu Ser His Phe Glu Val Lys Ser Gln Lys Tyr Asn Ala 100 105 <210> 257 <211> 1027 <212> DNA <213> Helicobacter pylori <220> <221> CDS <222> (51)...(974) <400> 257 ggcgcatgaa tgcattaaat gtaaagaacg cgtgttttta gaggaagata atg gct 56 Met Ala 1 aaa gaa aat ccg cct atc gtt ttt ggg cct gtt tta tcc agg cgt ttt 104 Lys Glu Asn Pro Pro Ile Val Phe Gly Pro Val Leu Ser Arg Arg Phe ggg aag tot ttg ggc gtg gat ota tcg ccc tot aaa aaa caa tgc aat 152 Gly Lys Ser Leu Gly Val Asp Leu Ser Pro Ser Lys Lys Gln Cys Asn 25 tac aat tgc att tat tgc gag ttg ggt aaa gcc aag ccc att gaa cgc 200 Tyr Asn Cys Ile Tyr Cys Glu Leu Gly Lys Ala Lys Pro Ile Glu Arg 35 40 atg gaa gaa gtg atc aaa gtg gaa acc ttg att aac gcc att caa aac 248 Met Glu Glu Val Ile Lys Val Glu Thr Leu Ile Asn Ala Ile Gln Asn 55 gcc cta aac aac ctc acc acc ccc att gat gtt tta acc att acc gct 296 Ala Leu Asn Asn Leu Thr Thr Pro Ile Asp Val Leu Thr Ile Thr Ala 70 75 80 aat ggc gaa ccc acg cta tac cct cat tta tta gag ctt atc caa agc 344 Asn Gly Glu Pro Thr Leu Tyr Pro His Leu Leu Glu Leu Ile Gln Ser 85 90 atc aag cct ttt tta aag ggc gtt aaa act ttg att tta agc aat ggc 392 Ile Lys Pro Phe Leu Lys Gly Val Lys Thr Leu Ile Leu Ser Asn Gly 105 110 tcg ctc ttt tat gag cca aaa gtc cag caa gcc tta aag gaa ttt gac 440 Ser Leu Phe Tyr Glu Pro Lys Val Gln Gln Ala Leu Lys Glu Phe Asp 115 120 125

Gln Ser Cys Gln His Gly Ile Ile Ala Asn Ser Thr Tyr Ser Trp Trp

	_					-	-				aaa Lys					488
	_						_			_	att Ile					536
_	_									-	gtg Val	-	_		_	584
		_				_	-				tta Leu 190				_	632
											gat Asp					680
-	-			-			-				agc Ser	-	-	~	_	728
											gtg Val					776
											tgc Cys			-	-	824
											gca Ala 270					872
											gaa Glu					920
											ttg Leu					968
gcg Ala		taac	ctco	at t	tgta	agtt	t ta	cctt	actt	tag	ggat	agc	ttaa	agctt	:tt	1024
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<211> 308

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<213> Helicobacter pylori

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                                            60
Gln Asn Ala Leu Asn Asn Leu Thr Thr Pro Ile Asp Val Leu Thr Ile
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                                   90
Gln Ser Ile Lys Pro Phe Leu Lys Gly Val Lys Thr Leu Ile Leu Ser
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Asn Gly Ser Leu Phe Tyr Glu Pro Lys Val Gln Gln Ala Leu Lys Glu
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                                        155
Gly Ile Leu Arg Phe Ser Gln Ile Tyr Gln Gly Gln Leu Val Ala Glu
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Thr Ile Asp Arg Pro Ser Ser Phe Lys Ala Pro Lys Leu Ser Glu Asp
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Glu Leu Lys Cys Ser Leu Phe Phe Glu Gly Leu Cys Val Ser Leu
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                                        235
Pro Lys Arg Ser Ile Thr Gln Ala Lys Lys Leu Ile Ser Cys Gly Ile
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Ala Pro Leu Ile Leu Asp Ser Asn Ala Phe Lys His Leu Glu Thr Leu
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					aaa Lys											305
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			_	_	aaa Lys 95	_										401
					gca Ala						-	-	_	-		449
-		_		_	ata Ile	-		-		_		_	_	_	_	497
					gcg Ala											545
					aag Lys						_		_			593
					aga Arg 175											641
	_	-			gag Glu		-	_				_				689
					att Ile											737

	caa Gln															785
	att Ile 235															833
	aaa Lys		_								_					881
	aac Asn								_	~ ~						929
	atc Ile															977
	gct Ala		_									-			-	1025
	tta Leu 315	_		_		-	_		-	-		-	_			1073
	gaa Glu															1121
	agg Arg	_				-			_							1169
	ctt Leu													-		1217
	gca Ala															1265
	agt Ser 395				tagg	gcata	ag a	aaat	aaga	aa ca	accat	aaaa	a tcg	;tttt	tag	1320
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170

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	_		_		aaa Lys					_					680
_	_				cga Arg	-		_	_						728
		-		_	gat Asp										776
					ctt Leu										824
				_	ggc Gly 265			-	-						872
					aaa Lys										920
	_				acg Thr		-			_				_	968
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		-		-	gcg Ala		-				_		_	_		536
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					ctc Leu 360			-		-	tagt	taaa	aaa (	gagat	taatc _.	1165
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							ggc Gly									440
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							ttg Leu									536
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	_	_			_	_					gtg Val			_		968
		_				_					gcc Ala	_		_		1016
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											ctc Leu					1208
											act Thr					1256
											agc Ser					1304
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ctt to Leu So 195																680
aag ga Lys G																728
ggt aa Gly Ly																776
tac go Tyr G	ly					_	_				-					824
ggc aa Gly As					_				_					-	_	872
ttg ca Leu Gi 275												taaa	aacco	ctt		918
agaaga	att	aa a	aggaa	aaga	ag at	gaaa	attt	taa	acgg	gatt	agc					961
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150
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145
Lys His Gly Ser Lys Gly Met Asp Tyr Leu Ala Gly Glu Met Pro Ala
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                165
Met Glu Leu Asp Glu Lys Asp Ala Lys Ala Ile Ala Ser Tyr Val Met
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Ala Glu Leu Ser Ser Val Lys Lys Thr Lys Asn Pro Gln Leu Ile Asp
                            200
                                                 205
Lys Gly Lys Glu Leu Phe Glu Ser Met Gly Cys Thr Gly Cys His Gly
                        215
                                             220
Asn Asp Gly Lys Gly Leu Gln Glu Asn Gln Val Phe Ala Ala Asp Leu
                                        235
                    230
Thr Ala Tyr Gly Thr Glu Asn Phe Leu Arg Asn Ile Leu Thr His Gly
                                     250
Lys Lys Gly Asn Ile Gly His Met Pro Ser Phe Lys Tyr Lys Asn Phe
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Ser Asp Leu Gln Val Lys Ala Leu Leu Asn Leu Ser Asn Arg
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                                                         Met Lys
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ttt tta aac gga tta gca ggg aat tta ctg att gtg gtt att tta ttg
                                                                   104
Phe Leu Asn Gly Leu Ala Gly Asn Leu Leu Ile Val Val Ile Leu Leu
                             10
tgt gtg gcc gtt ttt ttt acg ctc aaa gcg atc cat atc caa aaa gag
                                                                   152
Cys Val Ala Val Phe Phe Thr Leu Lys Ala Ile His Ile Gln Lys Glu
                         25
caa gcc acc aat tat tac cgc tat aag gat att aac gct tta gag aca
                                                                   200
Gln Ala Thr Asn Tyr Tyr Arg Tyr Lys Asp Ile Asn Ala Leu Glu Thr
                     40
                                         45
aaa aac acc caa aac cgg gct aac tat gaa tta gtc aat caa ggg agt
                                                                   248
Lys Asn Thr Gln Asn Arg Ala Asn Tyr Glu Leu Val Asn Gln Gly Ser
                 55
                                     60
aaa aaa tgaaattcac gactttagaa aaaattttag ctttgatggt agtagcgacc
                                                                   304
Lys Lys
att
                                                                   307
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                                 25
Lys Glu Gln Ala Thr Asn Tyr Tyr Arg Tyr Lys Asp Ile Asn Ala Leu
                             40
                                                 45
Glu Thr Lys Asn Thr Gln Asn Arg Ala Asn Tyr Glu Leu Val Asn Gln
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Gly Ser Lys Lys
65
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                                                         Met Gly
cga gcg ttt gaa tac aga aga gcg gct aaa gaa aaa cga tgg gat aag
                                                                    104
Arg Ala Phe Glu Tyr Arg Arg Ala Ala Lys Glu Lys Arg Trp Asp Lys
                              10
atg agt aag gtt ttc cca aag ctc gct aaa gcg atc act cta gcg gca
                                                                   152
Met Ser Lys Val Phe Pro Lys Leu Ala Lys Ala Ile Thr Leu Ala Ala
     20
aaa gat ggc ggg agc gaa ccg gac acg aac gcc aaa cta cga aca gcg
                                                                   200
Lys Asp Gly Gly Ser Glu Pro Asp Thr Asn Ala Lys Leu Arg Thr Ala
 35
                      40
att tta aac gct aaa gcg caa aac atg cct aaa gac aat att gac gca
                                                                   248
Ile Leu Asn Ala Lys Ala Gln Asn Met Pro Lys Asp Asn Ile Asp Ala
                 55
                                      60
gcg att aaa aga gcg agc agt aaa gaa ggg aat ttg agt gaa atc act
                                                                   296
Ala Ile Lys Arg Ala Ser Ser Lys Glu Gly Asn Leu Ser Glu Ile Thr
                                 75
tat gaa ggt aag gcg aat ttt ggc gtg cta atc atc atg gaa tgc atg
                                                                   344
Tyr Glu Gly Lys Ala Asn Phe Gly Val Leu Ile Ile Met Glu Cys Met
                             90
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act gat aac o Thr Asp Asn 1 100						s Ser				392
aaa acg caa ( Lys Thr Gln ( 115	Gly Ala			o Asn (						440
ttt aac cga a Phe Asn Arg I										488
tta aaa ctc a Leu Lys Leu S	_		_	u Phe	_		_			536
ttg gaa gaa t Leu Glu Glu I 165			-	_					_	584
tat aac agc t Tyr Asn Ser I 180						u Ser				632
ccc att tta a Pro Ile Leu I 195	Lys Ala			g Ile A						680
ttg aat gac ( Leu Asn Asp (										728
gaa gac gat g Glu Asp Asp A				u Tyr 1						770
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Ala Ala Lys A		Gly Ser			Thr As	n Ala 45		Leu	Arg	
Thr Ala Ile I	Leu Asn A	Ala Lys 55		n Asn N	Met Pr 60	o Lys	Asp	Asn	Ile	
Asp Ala Ala 1	Ile Lys A		Ser Se	r Lys (			Leu	Ser	Glu	

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Ile Thr Tyr Glu Gly Lys Ala Asn Phe Gly Val Leu Ile Ile Met Glu
                                    90
                85
Cys Met Thr Asp Asn Pro Thr Arg Thr Ile Ala Asn Leu Lys Ser Tyr
            100
                                105
                                                     110
Phe Asn Lys Thr Gln Gly Ala Ser Ile Val Pro Asn Gly Ser Leu Glu
                            120
                                                125
Phe Met Phe Asn Arg Lys Ser Val Phe Glu Cys Leu Lys Asn Glu Val
                        135
                                            140
Glu Asn Leu Lys Leu Ser Leu Glu Asp Leu Glu Phe Ala Leu Ile Asp
                    150
                                        155
Tyr Gly Leu Glu Glu Leu Glu Glu Val Glu Asp Lys Ile Ile Ile Arg
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Gly Asp Tyr Asn Ser Phe Lys Leu Leu Asn Glu Gly Phe Glu Ser Leu
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                                185
Lys Leu Pro Ile Leu Lys Ala Ser Leu Gln Arg Ile Ala Thr Thr Pro
                            200
Ile Glu Leu Asn Asp Glu Gln Met Glu Leu Thr Glu Lys Leu Leu Asp
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                                            220
Arg Ile Glu Asp Asp Asp Val Val Ala Leu Tyr Thr Asn Ile Glu
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aagcgataga atacaacaaa aagggtcagg ggcattttat aggcacc atg gtt gtc
                                                                   116
                                                 . Met Val Val
gct aag aat gaa gac aac aaa aaa ttg tat gac atc att gac ggc cag
                                                                   164
Ala Lys Asn Glu Asp Asn Lys Lys Leu Tyr Asp Ile Ile Asp Gly Gln
     5
                         10
caa cga acg act acc atc ttc atg ctc ttg cat gtc ttg gcg aac aaa
                                                                   212
Gln Arg Thr Thr Ile Phe Met Leu Leu His Val Leu Ala Asn Lys
 20
                     25
                                         30
caa aac gag aaa gac aag caa gaa aca aga aaa tat cta tac caa aag
                                                                   260
Gln Asn Glu Lys Asp Lys Gln Glu Thr Arg Lys Tyr Leu Tyr Gln Lys
                 40
ggg gaa tta aaa tta gaa gtc gcc ccc aaa aac caa agc ttc ttc aaa
                                                                   308
Gly Glu Leu Lys Leu Glu Val Ala Pro Lys Asn Gln Ser Phe Phe Lys
             55
                                 60
acg ctc ttg gaa gcg gca gaa aag gag aat atc agc cag aaa aag atg
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75

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Thr Leu Leu Glu Ala Ala Glu Lys Glu Asn Ile Ser Gln Lys Lys Met
         70
                             75
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cag aca ccg agg gca agc aaa atc ttt ttg aag ttt tgaaggctat
Gln Thr Pro Arg Ala Ser Lys Ile Phe Leu Lys Phe
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Asp Gly Gln Gln Arg Thr Thr Ile Phe Met Leu Leu His Val Leu
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Ala Asn Lys Gln Asn Glu Lys Asp Lys Gln Glu Thr Arg Lys Tyr Leu
Tyr Gln Lys Gly Glu Leu Lys Leu Glu Val Ala Pro Lys Asn Gln Ser
Phe Phe Lys Thr Leu Leu Glu Ala Ala Glu Lys Glu Asn Ile Ser Gln
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Lys Lys Met Gln Thr Pro Arg Ala Ser Lys Ile Phe Leu Lys Phe
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                                                           Met
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                                                                   104
Ser Asp Asn Glu Arg Thr Ile Val Val Arg Val Leu Lys Phe Asp Pro
caa agc gcg gtg agt aag ccg cat ttt aaa gag tat cag ttg aaa gaa
                                                                   152
Gln Ser Ala Val Ser Lys Pro His Phe Lys Glu Tyr Gln Leu Lys Glu
                             25
acg cca tcc atg acg ctt ttt atc gct ttg aac ctc att aga gag cat
                                                                   200
Thr Pro Ser Met Thr Leu Phe Ile Ala Leu Asn Leu Ile Arg Glu His
    35
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														att Ile		248
		_		_	_	_			-					tgt Cys 80		296
														ccc Pro		344
	-						-	_	_					gat Asp		392
		-	_						_					aaa Lys	_	440
														gaa Glu		488
														tgt Cys 160		536
														gct Ala	-	584
														gaa Glu		632
														gtt Val		680
										act Thr 220		cc t	aaag	yaatt	a	728
										tgt tag					gccgc	788 840
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Glu Thr Pro Ser Met Thr Leu Phe Ile Ala Leu Asn Leu Ile Arg Glu
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His Gln Asp Pro Asp Leu Ser Phe Asp Phe Val Cys Arg Ala Gly Ile
Cys Gly Ser Cys Ala Met Met Val Asn Gly Arg Pro Arg Leu Ala Cys
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                                        75
Lys Thr Leu Thr Ser Ser Phe Glu Ser Gly Val Ile Thr Leu Met Pro
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                                    90
Met Pro Ser Phe Thr Leu Ile Lys Asp Leu Ser Val Asn Thr Gly Asp
            100
                                105
Trp Phe Leu Asp Met Thr Lys Arg Val Glu Ser Trp Ala His Ser Lys
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Glu Glu Val Asp Ile Thr Arg Pro Glu Lys Arg Val Glu Pro Asp Glu
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                                            140
Ala Gln Glu Val Phe Glu Leu Asp Arg Cys Ile Glu Cys Gly Cys Cys
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Tie Ala Ser Cys Gly Thr Lys Leu Met Arg Pro Asn Phe Ile Gly Ala
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Ala Gly Met Asn Arg Ala Met Arg Phe Met Ile Asp Ser His Asp Glu
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Arg Asn Asp Asp Phe Tyr Glu Leu Val Gly Asp Asp Asp Gly Val
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Phe Gly Cys Met Ser Leu Ile Ala Cys His Asp Thr Cys Pro
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                                                                  106
Leu Leu Phe Ala Thr Pro Thr Ile Ser Pro Phe Leu Pro Phe Lys Asn
        5
ccc atg att ttc ctt tat aaa aat gaa atg att gtt tta aaa ttt tct
                                                                  154
Pro Met Ile Phe Leu Tyr Lys Asn Glu Met Ile Val Leu Lys Phe Ser
    20
                         25
aat too caa gac gog oto cog ato aac aag coa too acg ota toa ato
                                                                  202
Asn Ser Gln Asp Ala Leu Pro Ile Asn Lys Pro Ser Thr Leu Ser Ile
                     40
cct aaa att tct tta gcg ttt tgt gtg ttc acg ctc ccc cca tac aac
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Pro Gln Ser Ala Val Ser Lys Pro His Phe Lys Glu Tyr Gln Leu Lys

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Pro Lys Ile Ser Leu Ala Phe Cys Val Phe Thr Leu Pro Pro Tyr Asn
                                                          65
                 55
                                      60
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Lys Gly Val Phe
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<213> Helicobacter pylori
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Phe Ser Asn Ser Gln Asp Ala Leu Pro Ile Asn Lys Pro Ser Thr Leu
                            40
Ser Ile Pro Lys Ile Ser Leu Ala Phe Cys Val Phe Thr Leu Pro Pro
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Tyr Asn Lys Gly Val Phe
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gagagagag gtaaaactct tttatcaaat ccattgtttg gtggagagtt taaccccaga 180
gaatatagcc aaattagaag aaacgatcgc tccttttaga gctttttcta gcatagagtt 240
tttggatatt accgataaag aattagaacc acgccacaat tataataagc ttgatccttt 300
aatagcgagt gaaattaaaa aattgtattt aaaactcaat gctttttcgc aaaaacgctt 360
ttctaaa atg atc atg tgc cgt ttc ttt ttt gcc tcc ctt ttc ccc caa
        Met Ile Met Cys Arg Phe Phe Phe Ala Ser Leu Phe Pro Gln
tac gat aag atg atc atg ttt gat gtg gac act ttg ttt gtg aat gat
                                                                   457
Tyr Asp Lys Met Ile Met Phe Asp Val Asp Thr Leu Phe Val Asn Asp
15
                     20
                                         25
                                                              30
att agc gag agc ttt ttt atc ccc ctt gaa acg cat tat ttt ggg gct
                                                                   505
Ile Ser Glu Ser Phe Phe Ile Pro Leu Glu Thr His Tyr Phe Gly Ala
                 35
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								ata Ile 55							553
				_				gca Ala					_		601
_					_	-	_	caa Gln				-			649
	-				_			tta Leu				-			697
								ttc Phe							745
			_		_	-	_	tgt Cys 135			-	-	 		793
	_	-						gcc Ala			_		-	_	841
								gcg Ala							889
								gtc Val							937
		_			_			aaa Lys	_	_			_		985
								ctt Leu 215							1033
								gaa Glu				_	 -		1081
								cat His							1129
								gaa Glu							1177

caa aaa gtg aag cga ctg att aaa agg gtt ttt taaaaccctt tttaaactaa 1230 Gln Lys Val Lys Arg Leu Ile Lys Arg Val Phe 275 280

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Glu Ser Phe Phe Ile Pro Leu Glu Thr His Tyr Phe Gly Ala Val Arg
                            40
Glu Lys Asp Leu Ile Ala Ile Asn Arg Asn Ser Ala Lys Asp Leu Tyr
                        55
Glu Leu Arg Gln Met His Ala Lys Ser Ile Gly Ile Ala Asn Ala Phe
                    70
                                        75
Pro Asn Leu Glu Glu Ala Gln Ile Leu Phe Asp Asn Tyr Phe Asn Ala
                85
                                    90
Gly Phe Leu Ala Leu Asn Leu Lys Ser Trp Arg Lys Glu Asn Leu Glu
                                105
Asn Gln Leu Ile Thr Phe Phe Ile Leu Lys Asn Glu Lys Leu Leu Phe
                            120
                                                125
Asn Asp Gln Asp Ala Leu Cys Phe Val Cys Arg Gly Arg Ile Leu Glu
                        135
                                            140
Leu Pro Tyr Pro Tyr Asn Ala His Pro Ser Phe Leu Asp Thr Leu Ser
                   150
                                        155
Phe Pro Ser Ile Lys Glu Ala Arg Met Leu His Phe Trp Gly Asp Lys
                165
                                    170
Pro Trp Lys Leu Ser Val Ile Gly Ala Lys Lys Trp His Glu Ala
                                185
                                                    190
Leu Ile Gln Thr Pro Phe Lys Asp Ala Tyr Phe Asn Ala Ser Phe Leu
                            200
Asp His Leu Phe Glu Ser Leu Gln Asn Lys Asp Asn Glu Ile Lys Arg
                        215
                                            220
Arg Asp Glu Arg Ile Ile Glu Ala Leu Gln Ala Arg Asp Lys Ile Leu
                    230
                                        235
Ser Phe Ser Asp Lys Arg His Ser Phe Glu Ser Leu Leu Pro Lys Leu
                245
                                    250
Ser Ser Lys Leu Leu Ile Glu Phe Leu Leu Phe Lys Ala Lys Gln Lys
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Val Lys Arg Leu Ile Lys Arg Val Phe
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<212> DNA

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Ser Phe Ser Ala His Lys Phe His Gly Pro Lys Gly Ile Gly Gly Leu

	100					103					170					
					gtg Val 200											680
					cgc Arg											728
					atg Met											776
					aaa Lys											824
					atg Met											872
					agc Ser 280											920
					tct Ser											968
					gag Glu											1016
					cat His							_	_	_		1064
				Glu	att Ile	Asp	Lys			Glu		Phe				1112
gct Ala 355	gta Val	agg Arg	ttg Leu	aga Arg	aat Asn 360	att Ile	tca Ser	agc Ser	tct Ser	tat Tyr 365	taaa	aaga	at a	ıtaaa	aggaat	1165
caaa	atgg	jca a	aaca	atgat	t ta	agtgg	gttc	ggt	;							1198
<211 <212	)> 28 .> 36 !> PF !> He	55 RT	bact	er p	oylor	· ·i										
	)> 28															

190

180

Met Asp Pro Phe Leu Arg Asp His Tyr Gly Asn Pro Ser Ser Leu His

```
10
Gln Phe Gly Thr Glu Thr His Pro Ala Ile Ala Glu Ala Leu Asp Lys
Leu Tyr Lys Gly Ile Asn Ala Arg Asp Ile Asp Asp Val Ile Ile Thr
                            40
Ser Cys Ala Thr Glu Ser Asn Asn Trp Val Leu Lys Gly Val Tyr Phe
                        55
Asp Glu Cys Leu Lys Lys Gly Lys Asn His Ile Val Thr Thr Val Ala
                    70
                                        75
Glu His Pro Ala Val Arg Ser Thr Cys Asn Phe Leu Glu Ser Leu Gly
                                    90
                85
Val Glu Val Thr Tyr Leu Pro Ile Asn Glu His Gly Ser Ile Thr Ala
                               105
Glu Gln Val Lys Glu Ala Ile Thr Glu Lys Thr Ala Leu Val Ser Val
                            120
Met Trp Ala Asn Asn Glu Thr Gly Leu Ile Phe Pro Ile Glu Glu Ile
                        135
                                            140
Gly Ala Ile Cys Lys Glu Lys Gly Val Leu Phe His Thr Asp Ala Val
                   150
                                        155
Gln Ala Ile Gly Lys Ile Pro Val Asp Val Leu Lys Ala Asn Ala Asp
                                    170
Phe Leu Ser Phe Ser Ala His Lys Phe His Gly Pro Lys Gly Ile Gly
                                185
Gly Leu Tyr Ile Arg Ser Gly Val Gly Leu Thr Pro Leu Phe His Gly
        195
                           200
Gly Glu His Met Asn Gly Arg Arg Ser Gly Thr Leu Asn Val Pro Tyr
                       215
Ile Val Gly Met Gly Glu Ala Met Lys Leu Ala Val Glu His Leu Asp
                    230
                                        235
Tyr Glu Lys Glu Val Val Gly Lys Leu Arg Asp Lys Leu Glu Glu Ala
                245
                                    250
Leu Leu Lys Ile Pro Asp Val Met Val Val Gly Asp Arg Ile His Arg
                                265
Val Pro Asn Thr Thr Leu Val Ser Val Arg Gly Ile Glu Gly Glu Ala
                           280
Met Leu Trp Asp Leu Asn Arg Ser Asn Ile Ala Ala Ser Thr Gly Ser
                       295
                                            300
Ala Cys Ala Ser Glu Asp Leu Glu Ala Asn Pro Val Met Val Ala Ile
                                        315
                   310
Gly Ala Ser Lys Glu Leu Ala His Thr Ala Ile Arg Leu Ser Leu Ser
               325
                                   330
Arg Phe Asn Thr Glu Ala Glu Ile Asp Lys Thr Ile Glu Val Phe Ser
                                345
Gln Ala Ala Val Arg Leu Arg Asn Ile Ser Ser Ser Tyr
                            360
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<210> 285
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<211> 1273

<212> DNA

<213> Helicobacter pylori

<220>

<221> CDS

<222> (51)...(1220)

G	aa In	aag Lys 20	gat Asp	ttg Leu	ttg Leu	tct Ser	ggt Gly 25	ttt Phe	gtg Val	gtg Val	Gly	ctt Leu 30	tct Ser	gtg Val	atc Ile	cca Pro	152
Ğ															gtg Val		200
															ggg Gly 65		248
															att Ile		296
															gcg Ala		344
		_	_			_							_		ata Ile		392
A															gtg Val		440
															caa Gln 145		488
															atc Ile		536
														_	att Ile	_	584
	le														gcg Ala		632
															atc Ile		680

195	200	205	210
		ata atg ata gag ttg Ile Met Ile Glu Leu 220	
		acg ata gaa agc tta Thr Ile Glu Ser Leu 235	
_		gac ggc gtg agc gat Asp Gly Val Ser Asp 255	Lys Asn Lys
		aat atc atc tca ggg Asn Ile Ile Ser Gly 270	
		ggg cag tct atc att Gly Gln Ser Ile Ile 285	
		act ttt ttt gcc ggc Thr Phe Phe Ala Gly 300	
		gaa tat gtg gtt aag Glu Tyr Val Val Lys 315	•
		atg att tct ttc acc Met Ile Ser Phe Thr 335	Thr Phe Asn
		aaa atc aag ctc tat Lys Ile Lys Leu Tyr 350	
		gtt tta tac acg cat Val Leu Tyr Thr His 365	
		gtc aat gcg tta tgg Val Asn Ala Leu Trp 380	
aaa ggg att gca Lys Gly Ile Ala 390	tgaaatttta tttt	aaaaag ttgggtagct ag	ragatatgg 1260
ctccagatgt agg			1273
<210> 286 <211> 390 <212> PRT			

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<400> 286
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Asn Ile Gln Lys Asp Leu Leu Ser Gly Phe Val Val Gly Leu Ser Val
                                25
Ile Pro Glu Thr Ala Gly Phe Ala Ile Met Val Gly Leu Asp Val Gly
                            40
Val Ala Phe Tyr Thr Thr Phe Tyr Met Ala Phe Val Leu Ser Leu Phe
                        55
Gly Ala Arg Lys Ala Met Ile Ser Ala Ala Ala Gly Ser Val Ala Leu
Ile Leu Val Gly Val Val Lys Asn Tyr Gly Leu Glu Tyr Ala Gly Val
                                    90
                85
Ala Thr Leu Met Ala Gly Val Leu Gln Ile Leu Leu Gly Tyr Leu Lys
                                105
Ile Gly Asn Leu Leu Arg Phe Ile Pro Gln Ser Val Met Tyr Gly Phe
                           120
Val Asn Ala Leu Gly Ile Leu Leu Leu Met Glu Gln Phe Lys Phe Leu
                        135
                                            140
Gln Asn Gln Asn Leu Gly Val Phe Val Leu Leu Ala Ile Gly Ile Leu
                    150
                                        155
Ile Ile Tyr Leu Phe Pro Leu Ile Thr Lys Lys Ile Pro Ser Asn Leu
                165
                                    170
Ile Cys Ile Leu Ile Val Ser Ala Ile Ala Leu Ile Phe Asp Met His
                                185
Ala Pro Asn Leu Gly Ser Ile Glu Gln Gly Val Ser Gly Phe His Phe
                            200
                                                205
Ile Ile Ile Pro Lys Asn Leu Asp Phe Lys Ile Met Ile Glu Leu Leu
                        215
                                            220
Pro Tyr Ala Leu Ser Leu Ala Leu Val Gly Thr Ile Glu Ser Leu Leu
                   230
                                        235
Thr Ala Lys Thr Leu Asp Val Ile Leu Lys Asp Gly Val Ser Asp Lys
                245
                                    250
Asn Lys Glu Thr Lys Ala Gln Gly Leu Gly Asn Ile Ile Ser Gly Leu
                                265
Leu Gly Gly Met Thr Gly Cys Ala Leu Val Gly Gln Ser Ile Ile Asn
        275
                           280
                                                285
Ala Lys Ser Gly Ala Lys Thr Arg Leu Ser Thr Phe Phe Ala Gly Phe
                       295
Ser Leu Met Val Leu Ile Leu Val Phe Asn Glu Tyr Val Val Lys Ile
                    310
                                        315
Pro Ile Val Ala Val Val Ala Val Met Val Met Ile Ser Phe Thr Thr
                                    330
Phe Asn Phe Gln Ser Ile Ile Asn Ile Lys Lys Ile Lys Leu Tyr Asp
                                345
Thr Leu Asn Met Leu Leu Val Val Ala Val Leu Tyr Thr His Asn
                            360
Leu Ala Ile Gly Val Val Val Gly Val Leu Val Asn Ala Leu Trp Ile
                        375
Lys Ser Lys Gly Ile Ala
```

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<210> 287
<211> 1020
<212> DNA
<213> Helicobacter pylori
<220>
<221> CDS
<222> (386)...(964)
<400> 287
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gattggctct agggggagcg aattagcctt atggcaagcg aatcacatta aagaacgcct 120
gaaaaaagaa tgcttgatag aaagcgaaat tcaaatcgtt aagactaagg gcgataaaat 180
cttagacacc cctttaaata agatcggcgg taaggggcta ttcactaagg aattagaaga 240
attgctttta aagggcgcaa ttgatttggc ggtgcattct ttaaaagacg tgccggtcgt 300
gtttgaaaag gggttagact tggcatgcat caccaaaagg gctgatgtga gggacacttt 360
tttaagcgtg aaattccctg atttg atg agt ttg cct aaa ggg gca aag gtt
                            Met Ser Leu Pro Lys Gly Ala Lys Val
ggc acg act tct tta agg cgc tct atg cag atc aaa tta aag cgc cag
                                                                   460
Gly Thr Thr Ser Leu Arg Arg Ser Met Gln Ile Lys Leu Lys Arg Gln
                     15
gat ttg gac aca gaa agc ttg aga ggg aat gtc caa acc cgt ttg aaa
                                                                   508
Asp Leu Asp Thr Glu Ser Leu Arg Gly Asn Val Gln Thr Arg Leu Lys
                 3.0
aag ett gaa tge gga gaa ttt gae get ate att tta get gaa gee ggg
                                                                   556
Lys Leu Glu Cys Gly Glu Phe Asp Ala Ile Ile Leu Ala Glu Ala Gly
ttg tgc cgc cta gaa att caa gga gcg aaa tac cgc aag gct ttt agc
                                                                   604
Leu Cys Arg Leu Glu Ile Gln Gly Ala Lys Tyr Arg Lys Ala Phe Ser
                             65
gta gaa gaa atg att cet age atg ggt cag ggg get tta ggg gta gaa
                                                                   652
Val Glu Glu Met Ile Pro Ser Met Gly Gln Gly Ala Leu Gly Val Glu
     75
atg ctc aaa aac cac aag cat ttt gcc acg ctt caa aaa ctc aac gac
                                                                   700
Met Leu Lys Asn His Lys His Phe Ala Thr Leu Gln Lys Leu Asn Asp
 90
                     95
                                                             105
gag aaa agc gcg ttt tgc tgc cgt tta gaa agg gag ttt atc aag ggg
                                                                   748
Glu Lys Ser Ala Phe Cys Cys Arg Leu Glu Arg Glu Phe Ile Lys Gly
                110
                                    115
                                                         120
ctt aat gga ggg tgt cag atc cct ata ggc gtg cat gcg agt tta atg
                                                                   796
Leu Asn Gly Gly Cys Gln Ile Pro Ile Gly Val His Ala Ser Leu Met
            125
                                130
ggc gat agg gtt aaa atc cag gcg gtt tta ggc ttg cct aac ggg aaa
                                                                   844
Gly Asp Arg Val Lys Ile Gln Ala Val Leu Gly Leu Pro Asn Gly Lys
        140
                            145
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892
gaa gtc att act aaa gaa aaa caa ggg gat aaa act aaa gcg ttt gat
Glu Val Ile Thr Lys Glu Lys Gln Gly Asp Lys Thr Lys Ala Phe Asp
    155
                        160
tta gtt caa gag ctt tta gaa gaa ttt ttg caa agc ggg gct aaa gag
                                                                   940
Leu Val Gln Glu Leu Leu Glu Glu Phe Leu Gln Ser Gly Ala Lys Glu
                    175
                                        180
att tta gaa aag gcg cag ttg ttt taatgcgttt gtttatcgcg ctagttttgt
                                                                   994
Ile Leu Glu Lys Ala Gln Leu Phe
                190
                                                                   1020
tttggtggtg gttaagcttg aacgct
<210> 288
<211> 193
<212> PRT
<213> Helicobacter pylori
<400> 288
Met Ser Leu Pro Lys Gly Ala Lys Val Gly Thr Thr Ser Leu Arg Arg
                                    10
Ser Met Gln Ile Lys Leu Lys Arg Gln Asp Leu Asp Thr Glu Ser Leu
                                25
Arg Gly Asn Val Gln Thr Arg Leu Lys Lys Leu Glu Cys Gly Glu Phe
                            40
                                                 45
Asp Ala Ile Ile Leu Ala Glu Ala Gly Leu Cys Arg Leu Glu Ile Gln
                        55
Gly Ala Lys Tyr Arg Lys Ala Phe Ser Val Glu Glu Met Ile Pro Ser
                    70
                                        75
Met Gly Gln Gly Ala Leu Gly Val Glu Met Leu Lys Asn His Lys His
                                    90
Phe Ala Thr Leu Gln Lys Leu Asn Asp Glu Lys Ser Ala Phe Cys Cys
                                105
                                                     110
Arg Leu Glu Arg Glu Phe Ile Lys Gly Leu Asn Gly Gly Cys Gln Ile
        115
                            120
                                                125
Pro Ile Gly Val His Ala Ser Leu Met Gly Asp Arg Val Lys Ile Gln
                        135
                                            140
Ala Val Leu Gly Leu Pro Asn Gly Lys Glu Val Ile Thr Lys Glu Lys
                                        155
Gln Gly Asp Lys Thr Lys Ala Phe Asp Leu Val Gln Glu Leu Leu Glu
                165
                                    170
Glu Phe Leu Gln Ser Gly Ala Lys Glu Ile Leu Glu Lys Ala Gln Leu
                                185
Phe
<210> 289
<211> 265
<212> DNA
<213> Helicobacter pylori
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<220>

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<221> CDS
<222> (51)...(212)
<400> 289
taagcgcttt gcgtgtttaa aaattcctta tccaagctct cagcgtcttc atg aaa
                                                                   56
                                                         Met Lys
gct gta acc atc ttt cat gat aaa ttc cct cgc tct cac cag ccc aaa
                                                                   104
Ala Val Thr Ile Phe His Asp Lys Phe Pro Arg Ser His Gln Pro Lys
         5
tct tgg gcg gat ttc atc acg gaa ttt cgt gtg gat ttg ata gag atg
                                                                   152
Ser Trp Ala Asp Phe Ile Thr Glu Phe Arg Val Asp Leu Ile Glu Met
     20
                         25
                                              30
gac ggg caa ttg ctt gta act ttt aat gaa att agc ggc aat ttc ggt
                                                                   200
Asp Gly Gln Leu Leu Val Thr Phe Asn Glu Ile Ser Gly Asn Phe Gly
 35
                     40
                                                                   252
gat att ttc ttc taaagtgggg cttaaaacaa aatcattgtc ttttcggtct
Asp Ile Phe Phe
ttaaaaacca ata
                                                                   265
<210> 290
<211> 54
<212> PRT
<213> Helicobacter pylori
<400> 290
Met Lys Ala Val Thr Ile Phe His Asp Lys Phe Pro Arg Ser His Gln
                                    10
Pro Lys Ser Trp Ala Asp Phe Ile Thr Glu Phe Arg Val Asp Leu Ile
                                25
Glu Met Asp Gly Gln Leu Leu Val Thr Phe Asn Glu Ile Ser Gly Asn
                            40
                                                 45
Phe Gly Asp Ile Phe Phe
    50
<210> 291
<211> 1350
<212> DNA
<213> Helicobacter pylori
<220>
<221> CDS
<222> (73)...(1305)
<400> 291
ttttactttt tttttggtat tctaacaagc ttttaaataa tccaatctac tttgttttaa 60
ggataatatt tt atg gca gat gtc gtt gtg ggg atc cag tgg gga gat gag 111
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Met Ala Asp Val Val Gly Ile Gln Trp Gly Asp Glu

1 5 10

	-	-					agg Arg									159
							aat Asn									207
	_	_			-		tta Leu	_				-				255
							agc Ser									303
					_	-	ttt Phe 85		_		_			_		351
-	_	_	_	-			atc Ile	_				-			_	399
_			_				aac Asn			_						447
							atġ Met	-		_			_	_		495
							tta Leu			-			-			543
							aaa Lys 165				_					591
				-			ttt Phe				_				-	639
							agc Ser									687
							gly ggg									735
gat	tta	ggg	act	tac	cct	ttt	gta	aca	agc	tct	aac	acc	acg	agc	gct	783

Asp Leu G	Gly Thr 225	Tyr	Pro	Phe	Val	Thr 230	Ser	Ser	Asn	Thr	Thr 235	Ser	Ala	
agc gca t Ser Ala C		_												831
ata ggt a Ile Gly I 255			_				-							879
cct agc g Pro Ser G 270	_				_		_					-		927
gcg gag t Ala Glu F		_			_	_			-	_			-	975
gat ttg g Asp Leu V														1023
tta gcc t Leu Ala L 3	_			-	_		-			-			_	1071
gtg tgc g Val Cys V 335														1119
agc gat t Ser Asp L 350														1167
gaa aaa a Glu Lys S														1215
aga gag t Arg Glu T														1263
ctt att t Leu Ile S 4				_	_	_	-	_						1305
tgaaaaaat	t cgctt	ctgt	a tt	ggtg	rcaat	taa	aaac	cct	tgcg	rt				1350
<210> 292 <211> 411 <212> PRT <213> Hel		er p	ylor	·i										
<400> 292														

Met Ala Asp Val Val Val Gly Ile Gln Trp Gly Asp Glu Gly Lys Gly Lys Ile Val Asp Arg Ile Ala Lys Asp Tyr Asp Phe Val Val Arg Tyr 25 Gln Gly Gly His Asn Ala Gly His Thr Ile Val His Lys Gly Val Lys His Ser Leu His Leu Met Pro Ser Gly Val Leu Tyr Pro Lys Cys Lys 55 Asn Ile Ile Ser Ser Ala Val Val Val Ser Val Lys Asp Leu Cys Glu 70 Glu Ile Ser Ala Phe Glu Asp Leu Glu Asn Arg Leu Phe Val Ser Asp 85 90 Arg Ala His Val Ile Leu Pro Tyr His Ala Lys Lys Asp Ala Phe Lys 105 Glu Lys Ser Gln Asn Ile Gly Thr Thr Lys Lys Gly Ile Gly Pro Cys 120 Tyr Glu Asp Lys Met Ala Arg Ser Gly Ile Arg Met Gly Asp Leu Leu 135 140 Asp Asp Lys Ile Leu Glu Glu Lys Leu Asn Ala His Phe Lys Ala Ile 150 155 Glu Pro Phe Lys Lys Ala Tyr Asp Leu Gly Glu Asn Tyr Glu Lys Asp 165 170 Leu Met Gly Tyr Phe Lys Thr Tyr Ala Pro Lys Ile Cys Pro Phe Ile 185 Lys Asp Thr Thr Ser Met Leu Ile Glu Ala Asn Gln Lys Gly Glu Lys 195 200 205 Ile Leu Leu Glu Gly Ala Gln Gly Thr Leu Leu Asp Ile Asp Leu Gly 215 220 Thr Tyr Pro Phe Val Thr Ser Ser Asn Thr Thr Ser Ala Ser Ala Cys 230 235 Val Ser Thr Gly Leu Asn Pro Lys Ala Ile Asn Glu Val Ile Gly Ile 245 250 Thr Lys Ala Tyr Ser Thr Arg Val Gly Asn Gly Pro Phe Pro Ser Glu 265 Asp Thr Thr Pro Met Gly Asp His Leu Arg Thr Lys Gly Ala Glu Phe 280 285 Gly Thr Thr Thr Lys Arg Pro Arg Arg Cys Gly Trp Leu Asp Leu Val 295 300 Ala Leu Lys Tyr Ala Cys Ala Leu Asn Gly Cys Thr Gln Leu Ala Leu 310 315 Met Lys Leu Asp Val Leu Asp Gly Ile Asp Ala Ile Lys Val Cys Val 325 330 Ala Tyr Glu Arg Lys Gly Glu Arg Leu Glu Ile Phe Pro Ser Asp Leu 340 345 Lys Asp Cys Val Pro Ile Tyr Gln Thr Phe Lys Gly Trp Glu Lys Ser 360 Val Gly Val Arg Lys Leu Asp Asp Leu Glu Pro Asn Val Arg Glu Tyr 375 380 Ile Arg Phe Ile Glu Lys Glu Val Gly Val Lys Ile Arg Leu Ile Ser 390 Thr Ser Pro Glu Arg Glu Asp Thr Ile Phe Leu 405 410

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<211> 1363
<212> DNA
<213> Helicobacter pylori
<220>
<221> CDS
<222> (51)...(1310)
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                                                                    56
                                                         Val Asp
gta ttg agc gtg agc gaa atc aat gcg caa atc aaa gcc ctt tta gaa
                                                                    104
Val Leu Ser Val Ser Glu Ile Asn Ala Gln Ile Lys Ala Leu Leu Glu
                              10
                                                                    152
gcg act ttt ttg caa gtt agg gtt caa ggg gaa gtg agt aat ttg act
Ala Thr Phe Leu Gln Val Arg Val Gln Gly Glu Val Ser Asn Leu Thr
     20
                         25
atc cat aag gtg agc ggc cat gcg tat ttt tcg ctc aaa gac agc cag
                                                                    200
Ile His Lys Val Ser Gly His Ala Tyr Phe Ser Leu Lys Asp Ser Gln
                     40
tcg gtt att aaa tgc gtg ctg ttt aaa ggg aac gct aac agg ctc aaa
                                                                    248
Ser Val Ile Lys Cys Val Leu Phe Lys Gly Asn Ala Asn Arg Leu Lys
                                                          65
ttc gct tta aaa gaa ggg cag gaa gtg gtt gtt ttt ggg ggt att agc
                                                                    296
Phe Ala Leu Lys Glu Gly Gln Glu Val Val Phe Gly Gly Ile Ser
             70
gtg tat gtc cca agg ggg gat tat caa atc aat tgc ttt gaa ata gag
                                                                   344
Val Tyr Val Pro Arg Gly Asp Tyr Gln Ile Asn Cys Phe Glu Ile Glu
                             90
cct aag gat ata ggt tca tta act tta gct tta gag caa ttg aaa gaa
                                                                   392
Pro Lys Asp Ile Gly Ser Leu Thr Leu Ala Leu Glu Gln Leu Lys Glu
                        105
aaa tta cgc ctt aaa ggc tat ttt gat gaa gaa aat aaa tta ccc aaa
                                                                   440
Lys Leu Arg Leu Lys Gly Tyr Phe Asp Glu Glu Asn Lys Leu Pro Lys
115
                    120
                                         125
ccg cat ttt cct aaa cga gtg gca gtc atc act tct caa aat tca gcc
                                                                   488
Pro His Phe Pro Lys Arg Val Ala Val Ile Thr Ser Gln Asn Ser Ala
                135
                                     140
                                                         145
gct tgg gcg gac atg aaa aag atc gct tcc aaa cga tgg ccg atg tgt
                                                                   536
Ala Trp Ala Asp Met Lys Lys Ile Ala Ser Lys Arg Trp Pro Met Cys
            150
                                 155
                                                     160
gaa tta gtt tgt atc aac acc tta atg caa ggg gag ggc tgc gtt caa
                                                                   584
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Glu Leu Val Cys Ile Asn Thr Leu Met Gln Gly Glu Gly Cys Val Gln

agc	gtg	gtg	gaa	agc	atc	gtt	tat	gcg	gat	agt	ttt	cat	gac	aca	aaa
Ser	Val	Val	Glu	Ser	Ile	Val	Tyr	Ala	Asp	Ser	Phe	His	Asp	Thr	Lys
	180					185					190				

aac gct ttt gat gcg att gta gtg gct agg ggt ggg ggg agc atg gag Asn Ala Phe Asp Ala Ile Val Val Ala Arg Gly Gly Ser Met Glu gat ttg tat tct ttc aat gat gaa aaa atc gct gat gct ctg tat ttg Asp Leu Tyr Ser Phe Asn Asp Glu Lys Ile Ala Asp Ala Leu Tyr Leu gcc aaa acc ttc agc atg tca gct att ggg cat gag agc gat ttt tta Ala Lys Thr Phe Ser Met Ser Ala Ile Gly His Glu Ser Asp Phe Leu ttg age gat tta gtg gcg gat bta agg get tet aeg eet tea aac geg Leu Ser Asp Leu Val Ala Asp Leu Arg Ala Ser Thr Pro Ser Asn Ala atg gaa att tta ctc ccc agc agc gat gaa tgg ttg caa aga ctt gat Met Glu Ile Leu Leu Pro Ser Ser Asp Glu Trp Leu Gln Arg Leu Asp ggg ttt aat gtg aaa ttg cac cgc tcg ttt aaa act ttg ctc cac caa Gly Phe Asn Val Lys Leu His Arg Ser Phe Lys Thr Leu Leu His Gln aaa aag gcg cat tta gag cat tta gtg gct tct tta aaa cga ttg agt Lys Lys Ala His Leu Glu His Leu Val Ala Ser Leu Lys Arg Leu Ser ttt gaa aac aag cac cat tta aac gct tta aaa cta gaa aaa tta aaa Phe Glu Asn Lys His His Leu Asn Ala Leu Lys Leu Glu Lys Leu Lys atc gcc cta gaa aat aaa act cta gaa ttt tta cgc ttt aaa aaa acg Ile Ala Leu Glu Asn Lys Thr Leu Glu Phe Leu Arg Phe Lys Lys Thr Leu Leu Glu Lys Ile Ser Thr Gln Thr Leu Thr Ser Pro Phe Leu Gln act aaa aca gag cga ttg aac agg cta gaa aac gcc ctt aaa ctc gct Thr Lys Thr Glu Arg Leu Asn Arg Leu Glu Asn Ala Leu Lys Leu Ala cat gct aat ttg aaa tta ccc caa ttc ggg gcg ttg gtg agc aaa aat His Ala Asn Leu Lys Leu Pro Gln Phe Gly Ala Leu Val Ser Lys Asn 

aat caa gcg ata gaa tta gag gca tta aaa agg ggc gat aaa att gaa

```
Asn Gln Ala Ile Glu Leu Glu Ala Leu Lys Arg Gly Asp Lys Ile Glu
            390
                                395
                                                     400
tta agt aat gaa aaa acc aga gcg agc gct gaa att ttg agc gtg gat
                                                                   1304
Leu Ser Asn Glu Lys Thr Arg Ala Ser Ala Glu Ile Leu Ser Val Asp
agg gtg taggggtttg aaaaataata tttaaaacgc tatttgtttt atcttaaaat
                                                                   1360
Arg Val
    420
ttc
                                                                   1363
<210> 294
<211> 420
<212> PRT
<213> Helicobacter pylori
<400> 294
Val Asp Val Leu Ser Val Ser Glu Ile Asn Ala Gln Ile Lys Ala Leu
Leu Glu Ala Thr Phe Leu Gln Val Arg Val Gln Gly Glu Val Ser Asn
Leu Thr Ile His Lys Val Ser Gly His Ala Tyr Phe Ser Leu Lys Asp
                            40
                                                45
Ser Gln Ser Val Ile Lys Cys Val Leu Phe Lys Gly Asn Ala Asn Arg
                        55
Leu Lys Phe Ala Leu Lys Glu Gly Gln Glu Val Val Phe Gly Gly
                    70
Ile Ser Val Tyr Val Pro Arg Gly Asp Tyr Gln Ile Asn Cys Phe Glu
                                    90
Ile Glu Pro Lys Asp Ile Gly Ser Leu Thr Leu Ala Leu Glu Gln Leu
            100
                                105
Lys Glu Lys Leu Arg Leu Lys Gly Tyr Phe Asp Glu Glu Asn Lys Leu
                            120
                                                125
Pro Lys Pro His Phe Pro Lys Arg Val Ala Val Ile Thr Ser Gln Asn
                        135
                                            140
Ser Ala Ala Trp Ala Asp Met Lys Lys Ile Ala Ser Lys Arg Trp Pro
                    150
                                        155
Met Cys Glu Leu Val Cys Ile Asn Thr Leu Met Gln Gly Glu Gly Cys
                165
                                    170
Val Gln Ser Val Val Glu Ser Ile Val Tyr Ala Asp Ser Phe His Asp
                                185
Thr Lys Asn Ala Phe Asp Ala Ile Val Val Ala Arg Gly Gly Ser
                            200
Met Glu Asp Leu Tyr Ser Phe Asn Asp Glu Lys Ile Ala Asp Ala Leu
                        215
                                            220
Tyr Leu Ala Lys Thr Phe Ser Met Ser Ala Ile Gly His Glu Ser Asp
                   230
                                        235
Phe Leu Leu Ser Asp Leu Val Ala Asp Leu Arg Ala Ser Thr Pro Ser
                245
                                    250
Asn Ala Met Glu Ile Leu Leu Pro Ser Ser Asp Glu Trp Leu Gln Arg
                                265
                                                    270
Leu Asp Gly Phe Asn Val Lys Leu His Arg Ser Phe Lys Thr Leu Leu
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280

His Gln Lys Lys Ala His Leu Glu His Leu Val Ala Ser Leu Lys Arg 295 Leu Ser Phe Glu Asn Lys His His Leu Asn Ala Leu Lys Leu Glu Lys 305 310 315 Leu Lys Ile Ala Leu Glu Asn Lys Thr Leu Glu Phe Leu Arg Phe Lys 325 330 Lys Thr Leu Leu Glu Lys Ile Ser Thr Gln Thr Leu Thr Ser Pro Phe 340 345 350 Leu Gln Thr Lys Thr Glu Arg Leu Asn Arg Leu Glu Asn Ala Leu Lys 355 360 365 Leu Ala His Ala Asn Leu Lys Leu Pro Gln Phe Gly Ala Leu Val Ser 375 380 Lys Asn Asn Gln Ala Ile Glu Leu Glu Ala Leu Lys Arg Gly Asp Lys 395 390 Ile Glu Leu Ser Asn Glu Lys Thr Arg Ala Ser Ala Glu Ile Leu Ser 410 Val Asp Arg Val 420 <210> 295 <211> 1083 <212> DNA <213> Helicobacter pylori <220> <221> CDS <222> (51)...(1031) <400> 295 aaagacgcgc atggattctg ttgggtttga agaacctcaa aggggtttgg gtg ctt 56 Val Leu aag ggg tta aaa aaa gcg ttt aag gag agg ttt tgc tct caa gtg tat 104 Lys Gly Leu Lys Lys Ala Phe Lys Glu Arg Phe Cys Ser Gln Val Tyr atc tct ttt aat gtg gat cac aat ctt tta tcc act caa gtc ata agg 152 Ile Ser Phe Asn Val Asp His Asn Leu Leu Ser Thr Gln Val Ile Arg 20 atc aaa aac gat cgc att aaa gag aaa ttt ttt aaa act ttt gag act 200 Ile Lys Asn Asp Arg Ile Lys Glu Lys Phe Phe Lys Thr Phe Glu Thr 35 aaa gtg gag act aaa aat ggt gaa gtc cct att caa gcc tta aaa atc 248 Lys Val Glu Thr Lys Asn Gly Glu Val Pro Ile Gln Ala Leu Lys Ile 55 gcc aga act tat agc caa aaa tac ccc tac act tat ttt agc gcg atg 296 Ala Arg Thr Tyr Ser Gln Lys Tyr Pro Tyr Thr Tyr Phe Ser Ala Met 70 75 agt aaa gct aaa gag gtt tta tgc gaa aag cag gcg ttt gaa caa atc

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_	_			-		_	-			_	_			cgt Arg		440
		_	_		_		_		_			_		att Ile 145		488
_			_	_			_		_	_	_	_		ttg Leu		536
														gag Glu		584
								-	_			-		ttt Phe		632
			_	_	_		_		_	_			_	gtg Val	_	680
_		_	_	_			-	_		_	-		_	gaa Glu 225		728
Leu	Phe ggc	Leu	Ser	Glu 215 agc	Ile agt	Gln aag	Glu	Asp	Ile 220 agc	Asp	Ser	Leu ata	Glu	Glu	Ala	728
Leu ata Ile gct	Phe ggc Gly tat	Leu cta Leu	gac Asp 230	Glu 215 agc Ser	Ile agt Ser gaa	Gln aag Lys ggc	Glu gat Asp	Asp aat Asn 235 acg	Ile 220 agc Ser	Asp gaa Glu atc	Ser aaa Lys ccc	Leu ata Ile tta	Glu aca Thr 240 att	Glu 225 gag	Ala gac Asp	
ata Ile gct Ala	Phe ggc Gly tat Tyr	cta Leu agt Ser 245	gac Asp 230 ttg Leu	Glu 215 agc Ser att Ile	agt Ser gaa Glu	Gln aag Lys ggc Gly cgt	Glu gat Asp atg Met 250	Asp aat Asn 235 acg Thr	Ile 220 agc Ser aat Asn	Asp gaa Glu atc Ile	Ser aaa Lys ccc Pro	ata Ile tta Leu 255	Glu aca Thr 240 att Ile	Glu 225 gag Glu gca	Ala gac Asp gat Asp	776
Leu ata Ile gct Ala gtt Val	ggc Gly tat Tyr ttg Leu 260	cta Leu agt Ser 245 caa Gln	gac Asp 230 ttg Leu gag Glu	Glu 215 agc Ser att Ile gga Gly	agt Ser gaa Glu ttg Leu	Gln aag Lys ggc Gly cgt Arg 265 gtt	Glu gat Asp atg Met 250 ggc Gly	Asp aat Asn 235 acg Thr gtc Val	lle 220 agc Ser aat Asn tat Tyr	Asp gaa Glu atc Ile cat His	Ser aaa Lys ccc Pro tct Ser 270 caa	Leu ata Ile tta Leu 255 aga Arg	Glu aca Thr 240 att Ile gag Glu cac	Glu 225 gag Glu gca Ala	gac Asp gat Asp gac Asp	776 824

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Glu Lys His Ala Phe
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Glu Thr Lys Val Glu Thr Lys Asn Gly Glu Val Pro Ile Gln Ala Leu
Lys Ile Ala Arg Thr Tyr Ser Gln Lys Tyr Pro Tyr Thr Tyr Phe Ser
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Ala Met Ser Lys Ala Lys Glu Val Leu Cys Glu Lys Gln Ala Phe Glu
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Gln Ile Lys Gln Glu Asn Gln Asp Tyr His Ala Cys Glu Val Asn Gln
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Lys Tyr Cys Val Tyr Val Glu Ser Lys Asp Phe Leu Lys Asp Phe Lys
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                                                125
Arg Phe Lys Ile Gln Asp Val Asp Phe Leu Phe Ser Pro Phe Ser Leu
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Ile Tyr Asp Phe Val Arg Asp Asn Leu Glu Asn Lys Pro Leu Leu Tyr
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Leu Leu Glu Arg Ser Arg Phe Tyr Phe Leu Ile Ala Asp Lys Lys
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Glu Ile Phe Leu Ala Lys Ser Val Phe Leu Glu Glu Gln Pro Glu Glu
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Val Asp Leu Phe Leu Ser Glu Ile Gln Glu Asp Ile Asp Ser Leu Glu
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Glu Ala Ile Gly Leu Asp Ser Ser Lys Asp Asn Ser Glu Lys Ile Thr
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                                        235
Glu Asp Ala Tyr Ser Leu Ile Glu Gly Met Thr Asn Ile Pro Leu Ile
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Ala Asp Val Leu Gln Glu Gly Leu Arg Gly Val Tyr His Ser Arg Glu
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Ile Asp Phe Val Glu Lys Val Val Leu Asp Ser Cys Gln Ile His
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aat tta gcc cat tct caa gtg gca acg cct gtt ttt atg ccc gta ggc 156 Asn Leu Ala His Ser Gln Val Ala Thr Pro Val Phe Met Pro Val Gly 20 25 30

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Thr Gln Gly Cys Ile Lys Ser Leu Asp Ala Thr Asp Ala Gln Glu Ile
35 40 45

tta ggc gct aaa ctc att tta gcc aac acc tat cac atg tat tta agg 252 Leu Gly Ala Lys Leu Ile Leu Ala Asn Thr Tyr His Met Tyr Leu Arg 50 65

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caa ttt tat ggg agt ttt tta acc gat agt gga ggg ttt caa gcc ttt 348 Gln Phe Tyr Gly Ser Phe Leu Thr Asp Ser Gly Gly Phe Gln Ala Phe 85 90 95

agt ttg agc gat aat gtc aaa ttg caa gaa gat ggg atc gtt ttt aaa 396 Ser Leu Ser Asp Asn Val Lys Leu Gln Glu Asp Gly Ile Val Phe Lys 100 105 110

tcc cat att gat ggg agc aag cat cta ttc acg ccc gct aaa gtt ttg 444
Ser His Ile Asp Gly Ser Lys His Leu Phe Thr Pro Ala Lys Val Leu
115 120 125

gac att caa tat tct ttg aat agc gat att atg atg gtt tta gac gat 492 Asp Ile Gln Tyr Ser Leu Asn Ser Asp Ile Met Met Val Leu Asp Asp 130 135 140 145

											gaa Glu				540
											tac Tyr				588
	_	_	_					-			cag Gln	 	_		636
_		_	_	_		_				_	cat His 205	 		_	684
											agc Ser				732
	-						-		_		ccc Pro	-	_		780
											atc Ile				828
-	_			_	_		_	_		_	ccc Pro	_		_	876
		-				_					att Ile 285				924
											gaa Glu				972
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tct cat tgaatgatgg aatgcaaaaa tactaaaaaag cgttttttac catcaataaa

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Val	Gly	Lys	Thr	Arg 135	Ala	Phe	Ile	Lys	Ile 140	Gln	Glu	Gly	Cys	Asp 145	Phe	
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	-		_		att Ile				-	-			_			584
	-		-		gtt Val								-			632
	_	_		_	aat Asn 200			_	_					_	_	680
					cgc Arg											728
		-	_		tta Leu				-		-			-		776
					tta Leu											824
_		-	_		cgc Arg			-	-		-			-		872
					ttt Phe 280											920
					agc Ser											968
					cac His									_		1016
				_	ttg Leu	_		-	_		_	_	_	-		1064
					gcg Ala											1112

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                                        365
355
                                                                   1208
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Val Glu Val Gln Lys Asp Gly Glu Phe Lys Ala Leu Asp Gln Phe Phe
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                                    380
aac ccc att aaa atc aaa agc gat aag cct cta agg gct agt ttt tta
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Asn Pro Ile Lys Ile Lys Ser Asp Lys Pro Leu Arg Ala Ser Phe Leu
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Glu Ile Lys Glu Tyr Glu Ile Lys Glu Arg Glu Asn His Ala Val Phe
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Asn Gly Ala Asp Ser Ala Val Arg Ser Tyr Ala Lys Lys Met Ala Arg
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Lys Glu Leu Phe Glu Lys Gly Phe Leu Lys Gly Val Phe Gly His Asp
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Asn Lys Glu Lys Ile Asn Ala Leu Leu Gln Glu Lys Lys Arg Phe Phe
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Ile Asp Asp Asn Leu Glu Asn Lys His Leu Asp Thr Thr Met Val Ser
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Asp Phe Asp Cys Asn Tyr Cys Ile Ile Pro Ser Val Arg Gly Arg Ala
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Arg Ser Phe Glu Glu Arg Lys Ile Leu Glu Gln Val Gly Leu Leu Cys
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Ser Lys Gly Val Gln Glu Val Val Leu Thr Gly Thr Asn Val Gly Ser
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Tyr Gly Lys Asp Arg Gly Ser Asn Ile Ala Arg Leu Ile Lys Lys Leu
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                                                205
Ser Gln Ile Ala Gly Leu Lys Arg Ile Arg Ile Gly Ser Leu Glu Pro
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Asn Gln Ile Asn Asp Glu Phe Leu Glu Leu Leu Glu Glu Asp Phe Leu
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Glu Lys His Leu His Ile Ala Leu Gln His Ser His Asp Leu Met Leu
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Glu Thr Ile Ala Ser Lys Asn Phe Ala Ile Gly Thr Asp Phe Ile Val
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Gly His Pro Gly Glu Ser Gly Ser Val Phe Glu Lys Ala Phe Lys Asn
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                                             300
Leu Glu Ser Leu Pro Leu Thr His Ile His Pro Phe Ile Tyr Ser Lys
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                                        315
Arg Lys Asp Thr Pro Ser Ser Leu Met Thr Asp Ser Val Ser Leu Glu
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                                    330
                                                         335
Asp Ser Lys Lys Arg Leu Asn Ala Ile Lys Asp Leu Ile Phe His Lys
                                345
Asn Lys Ala Phe Arg Gln Leu Gln Leu Lys Leu Asn Thr Pro Leu Lys
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Ala Leu Val Glu Val Gln Lys Asp Gly Glu Phe Lys Ala Leu Asp Gln
                        375
Phe Phe Asn Pro Ile Lys Ile Lys Ser Asp Lys Pro Leu Arg Ala Ser
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Lys Phe Gly Leu Gly Val Tyr Leu Leu Leu Gly Ile Leu Gly Gly
tct ttg atc att cta gga gcg ata gtc gcg ccc att gtt ttc aaa gct
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Ser Leu Ile Ile Leu Gly Ala Ile Val Ala Pro Ile Val Phe Lys Ala
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tca age gtt tta cct gaa ttg cat ctg act ccc ttt gag age ggg aaa
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Ser Ser Val Leu Pro Glu Leu His Leu Thr Pro Phe Glu Ser Gly Lys
35
                     40
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Leu Met Ala Gln Ile Phe Val Arg Phe Asn Tyr Val Leu Gly Ala Ile
                 55
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														ttg Leu		344
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														cac His		440
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aat	gaga	gag t	tttt	ggct	ta c	catct	tagga	a aat	=							574
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Leu Phe Phe Glu Leu Glu Phe Leu Asp Thr Ala Ile Lys Ile Ile Thr

		165					170					175				
		Met				Ile									gat Aspʻ	•
Glu	Lys	Asn	Arg	Leu	Phe	Leu	Gln	aat Asn	Asp	Asp	Lys	Asn	Arg	Ser		(

aat gtg atg att gtg gat tta ttg cgt aac gat ttg agc cgc ttg gcc

632

680

728

Asn Val Met Ile Val Asp Leu Leu Arg Asn Asp Leu Ser Arg Leu Ala
215
220
225

tta aaa aat agc gtg aaa gtc aat caa ttg ttt gaa atc atc agc ttg 776 Leu Lys Asn Ser Val Lys Val Asn Gln Leu Phe Glu Ile Ile Ser Leu 230 235 240

cct age gtg tat caa atg ata age gag att gaa geg aaa ttg ccc cta 824
Pro Ser Val Tyr Gln Met Ile Ser Glu Ile Glu Ala Lys Leu Pro Leu
245 250 255

aaa acc agc ttg ttt gag att ttt aag gcg ttg ttc cct tgc ggc tct 872
Lys Thr Ser Leu Phe Glu Ile Phe Lys Ala Leu Phe Pro Cys Gly Ser
260 265 270

gtg acc gga tgc cct aaa atc aaa acc atg caa atc att gaa agt tta 920 Val Thr Gly Cys Pro Lys Ile Lys Thr Met Gln Ile Ile Glu Ser Leu 275 280 285 290

gaa aaa cgc cct agg ggg gtg tat tgc ggg gcg ata ggc atg gtt gaa 968 Glu Lys Arg Pro Arg Gly Val Tyr Cys Gly Ala Ile Gly Met Val Glu 295 300 305

gaa aaa aaa gcc ctt ttt agc gtg cct atc cgc act tta gaa aaa aga 1016 Glu Lys Lys Ala Leu Phe Ser Val Pro Ile Arg Thr Leu Glu Lys Arg 310 315 320

gtg cac gaa aat ttt ttg cat tta ggg gta ggg agt ggg gta act tat 1064 Val His Glu Asn Phe Leu His Leu Gly Val Gly Ser Gly Val Thr Tyr 325 330 335

aaa agt aaa gcg cca aaa gaa tat gaa gaa agc ttt ttg aaa tcc ttt 1112 Lys Ser Lys Ala Pro Lys Glu Tyr Glu Glu Ser Phe Leu Lys Ser Phe 340 345 350

ttt gtg atg ccc aaa ata gaa ttt gag att gta gaa acg atg aaa att 1160 Phe Val Met Pro Lys Ile Glu Phe Glu Ile Val Glu Thr Met Lys Ile 355 360 365 370

atc aaa aag gat caa aaa tta gag att aat aat aaa aac gcc cat aaa 1208 Ile Lys Lys Asp Gln Lys Leu Glu Ile Asn Asn Lys Asn Ala His Lys 375 380 385

gaa cgc tta atg aat agc act cga tat ttt aac ttt aaa tac gat gaa 1256

Glu Arg Leu Mo	et Asn Ser 90	Thr Arg Ty		Phe Lys	Tyr Asp Glu 400	
aat ctt tta ga Asn Leu Leu A 405	_	_				1304
ctc aat aaa a Leu Asn Lys L 420						1352
tta aaa agc c Leu Lys Ser L 435	-				_	1400
aat gat ttt t Asn Asp Phe L		_	_		_	1448
gct cga gcg c Ala Arg Ala L 4			l Met Phe			1496
aac cag gat to Asn Gln Asp Lo 485				-	_	1544
gaa atc cat aa Glu Ile His Aa 500						1592
aac ggg acg gg Asn Gly Thr G						1640
gca cct ttg ad Ala Pro Leu L	<del>-</del>				_	1688
att aac gcg c Ile Asn Ala Lo 5!			u Val Lys		taactataaa	1737
aacagagcgg cta	aaaacctc at	ttttagaa a	taggttacc	caa		1780
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1	5		10		15	
Ala Thr Asn Le		Leu Lys As 25	п мта гей	_	30	

Asn Arg Gly Asn Gly Tyr Phe Val Gly Tyr Leu Leu Tyr Glu Ala Arg Leu Ala Phe Leu Asp Glu Asn Phe Gln Ser Gln Thr Pro Phe Leu Tyr 55 Phe Glu Gln Phe Leu Glu Arg Lys Lys Tyr Ser Leu Glu Pro Leu Lys Glu His Ala Phe Tyr Pro Lys Ile His Ser Ser Leu Asp Gln Lys Thr 90 Tyr Phe Lys Gln Phe Lys Ala Val Lys Glu Arg Leu Lys Asn Gly Asp 105 Thr Tyr Gln Val Asn Leu Thr Met Asp Leu Phe Leu Asp Thr Lys Ala 120 Lys Pro Lys Arg Val Phe Lys Glu Val Val His Asn Gln Asn Thr Pro 135 140 Phe Lys Ala Phe Ile Glu Asn Glu Phe Gly Ser Val Leu Ser Phe Ser 150 155 Pro Glu Leu Phe Phe Glu Leu Glu Phe Leu Asp Thr Ala Ile Lys Ile 165 170 Ile Thr Lys Pro Met Lys Gly Thr Ile Ala Arg Ser Lys Asn Pro Leu 185 Ile Asp Glu Lys Asn Arg Leu Phe Leu Gln Asn Asp Asp Lys Asn Arg 200 205 Ser Glu Asn Val Met Ile Val Asp Leu Leu Arg Asn Asp Leu Ser Arg 215 220 Leu Ala Leu Lys Asn Ser Val Lys Val Asn Gln Leu Phe Glu Ile Ile 230 235 Ser Leu Pro Ser Val Tyr Gln Met Ile Ser Glu Ile Glu Ala Lys Leu 245 250 Pro Leu Lys Thr Ser Leu Phe Glu Ile Phe Lys Ala Leu Phe Pro Cys 265 Gly Ser Val Thr Gly Cys Pro Lys Ile Lys Thr Met Gln Ile Ile Glu 280 Ser Leu Glu Lys Arg Pro Arg Gly Val Tyr Cys Gly Ala Ile Gly Met 295 300 Val Glu Glu Lys Lys Ala Leu Phe Ser Val Pro Ile Arg Thr Leu Glu 310 315 Lys Arg Val His Glu Asn Phe Leu His Leu Gly Val Gly Ser Gly Val 330 325 Thr Tyr Lys Ser Lys Ala Pro Lys Glu Tyr Glu Glu Ser Phe Leu Lys 340 345 Ser Phe Phe Val Met Pro Lys Ile Glu Phe Glu Ile Val Glu Thr Met 360 Lys Ile Ile Lys Lys Asp Gln Lys Leu Glu Ile Asn Asn Lys Asn Ala 375 380 His Lys Glu Arg Leu Met Asn Ser Thr Arg Tyr Phe Asn Phe Lys Tyr 390 395 Asp Glu Asn Leu Leu Asp Phe Glu Leu Glu Lys Glu Gly Val Leu Arg 405 410 Val Leu Leu Asn Lys Lys Gly Lys Leu Ile Lys Glu Tyr Lys Thr Leu 425 Glu Pro Leu Lys Ser Leu Glu Ile Arg Leu Ser Glu Ala Pro Ile Asp 440 Lys Arg Asn Asp Phe Leu Tyr His Lys Thr Thr Tyr Ala Pro Phe Tyr 455 Gln Lys Ala Arg Ala Leu Ile Lys Lys Gly Val Met Phe Asp Glu Ile

Phe Tyr Asn Gln Asp Leu Glu Leu Thr Glu Gly Ala Arg Ser Asn Leu 485 490 Val Leu Glu Ile His Asn Arg Leu Leu Thr Pro Tyr Phe Ser Ala Gly 505 Ala Leu Asn Gly Thr Gly Val Val Gly Leu Leu Lys Lys Gly Leu Val 520 525 Gly His Ala Pro Leu Lys Leu Gln Asp Leu Gln Lys Ala Ser Lys Ile 535 Tyr Cys Ile Asn Ala Leu Tyr Gly Leu Val Glu Val Lys Ile Lys 550 555 <210> 305 <211> 958 <212> DNA <213> Helicobacter pylori <22û> <221> CDS <222> (51)...(905) <400> 305 ttagtggata ttttatacgc ttttattgat cctagaataa ggttgtcata atg gag 56 Met Glu 1 tot tit aga gag tit atc caa caa tic aaa aaa aat aag gca gcg gtc 104 Ser Phe Arg Glu Phe Ile Gln Gln Phe Lys Lys Asn Lys Ala Ala Val 5 10 gtt ggg gct tgg att gtg ctt tta ttg gta att tgc gcg att ttt gcg 152 Val Gly Ala Trp Ile Val Leu Leu Val Ile Cys Ala Ile Phe Ala 20 25 ccc ctt tta gcc ccg cat gat cct tat gtc caa aac gcg caa gat cgc 200 Pro Leu Leu Ala Pro His Asp Pro Tyr Val Gln Asn Ala Gln Asp Arg 40 ctt ttg aag cct ata tgg gag cat gga ggg aat gct aaa tac ctt tta 248 Leu Leu Lys Pro Ile Trp Glu His Gly Gly Asn Ala Lys Tyr Leu Leu ggc acc gat gat ttg ggg cgc gat att ttg agc cgc ttg atc tat ggg 296 Gly Thr Asp Asp Leu Gly Arg Asp Ile Leu Ser Arg Leu Ile Tyr Gly 70 gcc agg att tct tta acc ata ggg att gtt tct atg ggg att gcg gtg 344 Ala Arg Ile Ser Leu Thr Ile Gly Ile Val Ser Met Gly Ile Ala Val 85 90 95 ttt ttt ggc acg ata cta ggg cta ata gcg ggg tat ttt ggg ggg aaa 392 Phe Phe Gly Thr Ile Leu Gly Leu Ile Ala Gly Tyr Phe Gly Gly Lys 100 105

475

465

		_	_					atc Ile									440
								gtg Val									488
			_	_				Gly ggg									536
								cta Leu 170									584
								tct Ser									632
						_		atc Ile				_		_		_	680
								gaa Glu									728
								ccc Pro									776
								gct Ala 250									824
								atg Met				_	_				872
								aaa Lys				taaa	ıagga	igc t	tgca	atgatt	925
	ttag	gaagt	ta a	agat	ttaa	ıa aa	ctta	attt	tto	;							958
<210> 306 <211> 285 <212> PRT <213> Helicobacter pylori																	
		)> 30		Dho	Δτα	Gl.v	Dho	Tla	Cln	Cl n	Dho	Tura	Tvo	λ α~	Tura	ת ה	
	1				5			Ile Val		10					15		
	ATO	var	v a r	GTA	лта	тър	TIG	vaı	ח∈ת	neu	пеп	val	TTE	СУБ	чта	TIG	

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25
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Phe Ala Pro Leu Leu Ala Pro His Asp Pro Tyr Val Gln Asn Ala Gln
                            40
Asp Arg Leu Leu Lys Pro Ile Trp Glu His Gly Gly Asn Ala Lys Tyr
Leu Leu Gly Thr Asp Asp Leu Gly Arg Asp Ile Leu Ser Arg Leu Ile
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                    70
Tyr Gly Ala Arg Ile Ser Leu Thr Ile Gly Ile Val Ser Met Gly Ile
                85
                                    90
Ala Val Phe Phe Gly Thr Ile Leu Gly Leu Ile Ala Gly Tyr Phe Gly
                                105
Gly Lys Thr Asp Ala Ile Ile Met Arg Ile Met Asp Ile Met Phe Ala
                            120
                                                 125
Leu Pro Ser Ile Leu Leu Ile Val Ile Val Val Ala Val Leu Gly Pro
                        135
Ser Leu Thr Asn Ala Met Leu Ala Ile Gly Phe Val Gly Ile Pro Gly
145
                    150
                                         155
Phe Ala Arg Leu Val Arg Ser Ser Val Leu Gly Glu Lys Glu Lys Glu
                                   170
Tyr Val Ile Ala Ser Lys Ile Asn Gly Ser Ser His Leu Arg Leu Met
                                185
                                                     190
Cys Lys Val Ile Phe Pro Asn Cys Ile Ile Pro Leu Ile Val Gln Thr
                            200
                                                 205
Thr Met Gly Phe Ala Ser Thr Val Leu Glu Ala Ala Ala Leu Ser Phe
                        215
                                             220
Leu Gly Leu Gly Ala Gln Pro Pro Lys Pro Glu Trp Gly Ala Met Leu
                    230
                                         235
Met Asn Ser Met Gln Tyr Ile Ala Thr Ala Pro Trp Met Leu Val Phe
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                                    250
Pro Gly Val Met Ile Phe Leu Thr Val Met Ser Phe Asn Leu Val Gly
                                265
Asp Gly Ile Met Asp Ala Leu Asp Pro Lys Arg Thr Ser
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aaagcgcagc ncttcttgga caaaaaattt gtat atg gga gtc aat tac caa aca 115
                                      Met Gly Val Asn Tyr Gln Thr
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Gly	Ser	Ile 10	Asn	Leu	Met	Thr	Asn 15	Ile	His	Glu	Val	Arg 20	Glu	Val	Thr	
	tat Tyr 25															211
	aag Lys															259
	aac Asn					_	_			_			_			307
_	ttt Phe		_						_	-						355
	ggc Gly															403
	aat Asn 105												-	-		451
	ggc Gly											_				499
	agt Ser													-	-	547
	aac Asn															595
	gta Val															643
	aac Asn 185															691
	ttt Phe											737				
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His Glu Val Arg Glu Val Thr Asn Tyr Gln Thr Gly Tyr Thr Asn Ile
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Ile Thr Ser Val Asn Ser Val Lys Lys Leu Thr Asn Met Gly Ser Asn
                            40
Gly Ile Gly Leu Val Met Gly Tyr Asn His Phe Phe His Pro Asp Lys
                        55
                                            60
Ile Leu Gly Leu Arg Tyr Phe Ala Phe Leu Asp Trp Gln Gly Tyr Gly
                    70
Met Arg Tyr Pro Lys Gly Tyr Tyr Gly Gly Asn Asn Met Ile Thr Tyr
                85
                                    90
Gly Val Gly Val Asp Ala Val Trp Asn Phe Phe Gln Gly Ser Phe Tyr
                                105
Gln Asp Asp Ile Ser Val Asp Ile Gly Val Phe Gly Gly Ile Ala Ile
                           120
Ala Gly Asn Ser Trp Tyr Ile Gly Ser Lys Gly Gln Glu Leu Leu Gly
                       135
                                            140
Ile Thr Asn Ser Ser Ala Val Asp Asn Thr Ser Phe Gln Phe Leu Phe
                    150
                                        155
Asn Phe Gly Leu Lys Ala Leu Phe Val Asp Glu His Glu Phe Glu Ile
               165
                                   170
Gly Phe Lys Phe Pro Thr Ile Asn Asn Lys Tyr Tyr Thr Thr Asp Ala
                                185
Leu Lys Val Gln Met Arg Arg Val Phe Ala Phe Tyr Val Gly Tyr Asn
        195
                            200
Tyr His Phe
    210
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<211> 517
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<220>
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<222> (51)...(464)
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                                                        Met Val
                                                         1
ggg ggt gga acg gta aaa aaa gac ttg aag aaa gcc att caa tac tat
                                                                   104
Gly Gly Gly Thr Val Lys Lys Asp Leu Lys Lys Ala Ile Gln Tyr Tyr
                             10
gtt aaa gcg tgt gaa ttg aat gaa atg ttt ggg tgt ctg tca tta gtt
                                                                   152
Val Lys Ala Cys Glu Leu Asn Glu Met Phe Gly Cys Leu Ser Leu Val
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					aac Asn 40											200
					agt Ser											248
					tat Tyr											296
				-	tgt Cys				_		-					344
					tat Tyr											392
					ttt Phe 120											440
					aac Asn			taga	atttg	gaa a	ataaa	atgct	g tt	tttt	agct	494
ggct	ttca	atg t	tttt	gtaa	ac co	CC ·										517
<210 <211 <212	)> 31 l> 13 !> PF	10 88 RT		_	ac co											517
<210 <211 <212 <213	)> 31 l> 13 2> PF B> He	10 88 RT elico	obact	er p	oylo:	ri										517
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<210 <211 <212 <213 <400 Met 1 Tyr Leu Ser Asp 65	)> 31  > 13  > PP    32  Val   Tyr   Val   Lys   50   Phe	No. 10 Ser 35 Ala	Gly Lys 20 Asn Cys Glu	Gly 5 Ala Ser Glu Asn	Thr Cys Gln Leu Gly	Val Glu Ile Asn 55 Lys	Leu Asn 40 Ser Tyr	Asn 25 Lys Gly Val	10 Glu Gln Asn Lys	Met Lys Gly Lys 75	Phe Leu Cys 60 Asp	Gly Phe 45 Arg Leu	Cys 30 Gln Phe Arg	15 Leu Tyr Leu Lys	Ser Leu Gly Ala 80	517
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<210 <211 <212 <213 <400 Met 1 Tyr Leu Ser Asp 65 Ala Leu	)> 31  > 13  > PR    32  > 31  Val   Tyr  Val   Lys   50   Phe   Gln   Ile	Note that the second se	Gly Lys 20 Asn Cys Glu Tyr Gly 100	Gly 5 Ala Ser Glu Asn Ser 85 Tyr	Thr Cys Gln Leu Gly 70 Lys	Val Glu Ile Asn 55 Lys Ala Gln	Leu Asn 40 Ser Tyr Cys	Asn 25 Lys Gly Val Gly Ala 105	10 Glu Gln Asn Lys Leu 90 Gly	Met Lys Gly Lys 75 Asn	Phe Leu Cys 60 Asp Asp Gly	Gly Phe 45 Arg Leu Gln Val	Cys 30 Gln Phe Arg Asp Val 110	15 Leu Tyr Leu Lys Gly 95 Lys	Ser Leu Gly Ala 80 Cys Asn	517

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                                                         Val Asp
tca gag ggg ttt tcg cct tct att tat acc gac aag aca ggg cat ccc
                                                                   104
Ser Glu Gly Phe Ser Pro Ser Ile Tyr Thr Asp Lys Thr Gly His Pro
                             10
acg att ggc tat ggc tat aat ttg agc gtt tat tct tat gag ggt aag
                                                                   152
Thr Ile Gly Tyr Gly Tyr Asn Leu Ser Val Tyr Ser Tyr Glu Gly Lys
     20
                         25
cgt atc acc aaa aca tat ggg ctt tta act gac ata ctc tct tat ggg
                                                                   200
Arg Ile Thr Lys Thr Tyr Gly Leu Leu Thr Asp Ile Leu Ser Tyr Gly
35
                     40
                                          45
tgg tat aaa aat ttg gac gca atg agg aga atg gtc atc ttg gat ttg
                                                                   248
Trp Tyr Lys Asn Leu Asp Ala Met Arg Arg Met Val Ile Leu Asp Leu
                 55
                                      60
                                                          65
age tae aat tta gge ttg aac gga ctg etc aaa tte aag caa tte ate
                                                                   296
Ser Tyr Asn Leu Gly Leu Asn Gly Leu Leu Lys Phe Lys Gln Phe Ile
             70
aag gcc ata gag gat aaa aat tat gct ttg gct gtg gag aga ctg caa
                                                                   344
Lys Ala Ile Glu Asp Lys Asn Tyr Ala Leu Ala Val Glu Arg Leu Gln
aaa agc ccg tat ttc aat caa gtg aaa aaa gag cgt caa gga ata tgg
                                                                   392
Lys Ser Pro Tyr Phe Asn Gln Val Lys Lys Glu Arg Gln Gly Ile Trp
aaa ttt tgaaattgga gggttgcgaa aaacattgta agaaaaaata cgcaatagaa
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Lys Phe
115
aag
                                                                   451
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<400> 312

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His Pro Thr Ile Gly Tyr Gly Tyr Asn Leu Ser Val Tyr Ser Tyr Glu
                                25
Gly Lys Arg Ile Thr Lys Thr Tyr Gly Leu Leu Thr Asp Ile Leu Ser
                            40
Tyr Gly Trp Tyr Lys Asn Leu Asp Ala Met Arg Arg Met Val Ile Leu
                        55
Asp Leu Ser Tyr Asn Leu Gly Leu Asn Gly Leu Leu Lys Phe Lys Gln
                    70
                                        75
Phe Ile Lys Ala Ile Glu Asp Lys Asn Tyr Ala Leu Ala Val Glu Arg
                85
                                   90
Leu Gln Lys Ser Pro Tyr Phe Asn Gln Val Lys Lys Glu Arg Gln Gly
                                105
Ile Trp Lys Phe
        115
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                                                       Asn Pro Gln
age ttt ttc agg aat acg cca cat ttt ttg atc agc cga att ttt atc
                                                                   106
Ser Phe Phe Arg Asn Thr Pro His Phe Leu Ile Ser Arg Ile Phe Ile
     5
                         10
cac cag cat gag aat aaa aat cag aat att gac tat gag caa cca aac
                                                                   154
His Gln His Glu Asn Lys Asn Gln Asn Ile Asp Tyr Glu Gln Pro Asn
 20
                     25
gat aga agc aaa ttc cac gct cac cct ttc aag agc gtt tta aca acc
                                                                   202
Asp Arg Ser Lys Phe His Ala His Pro Phe Lys Ser Val Leu Thr Thr
                 40
caa acg cta cca ctt ggt ttt tta gag aga gaa aga gag aga aag caa
                                                                   250
Gln Thr Leu Pro Leu Gly Phe Leu Glu Arg Glu Arg Glu Arg Lys Gln
             55
aat ttt aag att gat tct caa atc tat tcc ttt gca aaa gtt aag att
                                                                   298
Asn Phe Lys Ile Asp Ser Gln Ile Tyr Ser Phe Ala Lys Val Lys Ile
                             75
ggg tgt ttt aac atg att ttt ggc ctg ctc gca tca agc cct tat ttt
                                                                   346
Gly Cys Phe Asn Met Ile Phe Gly Leu Leu Ala Ser Ser Pro Tyr Phe
                         90
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<213> Helicobacter pylori
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Ile Phe Ile His Gln His Glu Asn Lys Asn Gln Asn Ile Asp Tyr Glu
                                25
Gln Pro Asn Asp Arg Ser Lys Phe His Ala His Pro Phe Lys Ser Val
                            40
Leu Thr Thr Gln Thr Leu Pro Leu Gly Phe Leu Glu Arg Glu Arg Glu
Arg Lys Gln Asn Phe Lys Ile Asp Ser Gln Ile Tyr Ser Phe Ala Lys
                    7û
                                        75
Val Lys Ile Gly Cys Phe Asn Met Ile Phe Gly Leu Leu Ala Ser Ser
                                    90
Pro Tyr Phe
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                                                         Met Ser
tat ttt ttt aaa atc att ctg ggc aca agc gtg atc gtg ggg gtg ttg
                                                                   104
Tyr Phe Phe Lys Ile Ile Leu Gly Thr Ser Val Ile Val Gly Val Leu
ttg ggc ttg tgg cgt ttg act tac gat aag ttc tat ttc tcg ttg gtc
                                                                   152
Leu Gly Leu Trp Arg Leu Thr Tyr Asp Lys Phe Tyr Phe Ser Leu Val
                         25
ttt gtg ttg ctg ata cta ggg att gtc gct tgt agc tat att tct tta
                                                                   200
Phe Val Leu Ile Leu Gly Ile Val Ala Cys Ser Tyr Ile Ser Leu
35
                     40
                                         45
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aaa atg cat cag agg aaa tgc ttc gcc aag tgt ttc gtg aat agt gaa Lys Met His Gln Arg Lys Cys Phe Ala Lys Cys Phe Val Asn Ser Glu

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tct ttt tta tcc aag atg tta cac tcc cca ata atg gta att tgc ttt
                                                                   296
Ser Phe Leu Ser Lys Met Leu His Ser Pro Ile Met Val Ile Cys Phe
                                  75
                                                                   344
tat ttt att ttt tca att ttc aca tcc ata tcc atc gtc tat agc gtg
Tyr Phe Ile Phe Ser Ile Phe Thr Ser Ile Ser Ile Val Tyr Ser Val
                             90
         85
ctg gac tat gat cag atg atg tgg ggg ttt gtt ttt tgc act atc gtt
                                                                   392
Leu Asp Tyr Asp Gln Met Met Trp Gly Phe Val Phe Cys Thr Ile Val
    100
                        105
                                             110
gtt tgc gct gtg gtg ttt ggc acg ctt gaa aaa aat gct caa gag tac
                                                                   440
Val Cys Ala Val Val Phe Gly Thr Leu Glu Lys Asn Ala Gln Glu Tyr
115
                    120
                                         125
                                                             130
cat caa aga aga tta ttt gat gct gat gtc tagagaagtg agtgcttttg
                                                                   490
His Gln Arg Arg Leu Phe Asp Ala Asp Val
                135
                                                                   522
tggggactct tttcttcatt ggcttgagtt gc
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<212> PRT
<213> Helicobacter pylori
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Val Leu Leu Gly Leu Trp Arg Leu Thr Tyr Asp Lys Phe Tyr Phe Ser
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Leu Val Phe Val Leu Leu Ile Leu Gly Ile Val Ala Cys Ser Tyr Ile
Ser Leu Lys Met His Gln Arg Lys Cys Phe Ala Lys Cys Phe Val Asn
                        55
                                             60
Ser Glu Ser Phe Leu Ser Lys Met Leu His Ser Pro Ile Met Val Ile
                    70
                                        75
Cys Phe Tyr Phe Ile Phe Ser Ile Phe Thr Ser Ile Ser Ile Val Tyr
                85
                                    90
Ser Val Leu Asp Tyr Asp Gln Met Met Trp Gly Phe Val Phe Cys Thr
                                105
Ile Val Val Cys Ala Val Val Phe Gly Thr Leu Glu Lys Asn Ala Gln
                            120
Glu Tyr His Gln Arg Arg Leu Phe Asp Ala Asp Val
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                        135
                                             140
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170

Val Gly Arg Ala 180		l Gln Leu			
gaa tgc gct ctc Glu Cys Ala Leu 195					
tta agc gcg atc Leu Ser Ala Ile		Tyr Gln	_		
aat tac aag ctt Asn Tyr Lys Leu 230					
gaa caa aat att Glu Gln Asn Ile 245					
ttt atc atg cat Phe Ile Met His 260		Ala Leu		Ser Leu Gly	
gac aca gag agt Asp Thr Glu Ser 275					
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Ile Asp Arg Ala Gly Ile Val Gly Glu Asp Gly Glu Thr His Gln Gly
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Leu Leu Asp Val Ser Tyr Leu Arg Ser Ile Pro Asn Met Val Ile Phe
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Ala Pro Arg Asp Asn Glu Thr Leu Lys Asn Ala Val Arg Phe Ala Asn
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Glu His Asp Ser Ser Pro Cys Ala Phe Arg Tyr Pro Arg Gly Ser Phe
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Ala Leu Lys Glu Gly Val Phe Glu Pro Ser Gly Phe Val Leu Gly Gln
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Ser Glu Leu Leu Lys Lys Glu Gly Glu Ile Leu Leu Ile Gly Tyr Gly
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                                    170
Asn Gly Val Gly Arg Ala His Leu Val Gln Leu Ala Leu Lys Glu Lys
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Asn Ile Glu Cys Ala Leu Leu Asp Leu Arg Phe Leu Lys Pro Leu Asp
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Pro Asn Leu Ser Ala Ile Val Ala Pro Tyr Gln Lys Leu Tyr Val Phe
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Ser Asp Asn Tyr Lys Leu Gly Gly Val Ala Ser Ala Ile Leu Glu Phe
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Leu Ser Glu Gln Asn Ile Leu Lys Pro Val Lys Ser Phe Glu Ile Ile
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Asp Glu Phe Ile Met His Gly Asn Thr Ala Leu Val Glu Lys Ser Leu
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                                                                   104
Leu Leu Phe Thr Gly Ala Cys Gly Tyr Ile Gly Ser His Thr Ala Arg
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gcg ttt tta gaa aaa acc aaa gaa aat atc att att gta gat gac tta
                                                                   152
Ala Phe Leu Glu Lys Thr Lys Glu Asn Ile Ile Ile Val Asp Asp Leu
                         25
age ace ggt ttt tta gag cac ctc aaa gcg tta gag cat tat tac cct
                                                                   200
Ser Thr Gly Phe Leu Glu His Leu Lys Ala Leu Glu His Tyr Tyr Pro
 35
                     40
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		~	-	ttt Phe 55					_	•	-	_			248
-	-		_	aat Asn	_	_	_			_			_	_	296
_				gct Ala					_	_		_		-	344
	-			acc Thr			_							_	392
				cat His											440
				gaa Glu 135				_	_		_	_	_		488
				cct Pro									-		536
				tct Ser											584
				gct Ala		-	_	_			-				632
				caa Gln											680
				gcg Ala 215							-				 728
				aca Thr											776
				gct Ala											824
				agc Ser											872

260 265 270	
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att gcc aat aac gct aaa atc tta caa aac acc tct ttc aaa cc Ile Ala Asn Asn Ala Lys Ile Leu Gln Asn Thr Ser Phe Lys Pr 310 315 320	
tat aac aac cta gac acc att atc aaa agc gct cta gat tgg ga Tyr Asn Asn Leu Asp Thr Ile Ile Lys Ser Ala Leu Asp Trp Gl 325 330 335	
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Leu Arg Tyr Phe Asn Val Ala Gly Ala Cys Met His Asn Asp Tyr Thr

Thr Pro Tyr Thr Leu Gly Gln Arg Thr Leu Asn Ala Thr His Leu Ile

195					200					205				
Lys Ile Ala 210	Сув	Glu	Cys	Ala 215	Val	Gly	Lys	Arg	Lys 220	Lys	Met	Gly	Ile	
Phe Gly Thr 225	Asn	Tyr	Pro 230	Thr	Arg	Asp	Gly	Thr 235	Cys	Ile	Arg	Asp	Tyr 240	
Ile His Val	_	Asp 245		Ala	Asn	Ala	His 250	Leu	Ala	Ser	Tyr	Gln 255	Thr	
Leu Leu Glu			Lys	Ser	Glu	Ile 265		Asn	Val	Gly	Tyr 270	Asn	Gln	
Gly His Ser		Lys	Glu	Val	Ile 280	Glu	Lys	Val	Lys	Glu 285	Ile	Ser	Asn	
Asn Asp Phe	Leu	Val	Glu	Ile 295	Leu	Asp	Lys	Arg	Gln 300	Gly	Asp	Pro	Ala	
Ser Leu Ile 305	Ala	Asn	Asn 310	Ala	Lys	Ile	Leu	Gln 315	Asn	Thr	Ser	Phe	Lys 320	
Pro Leu Tyr		Asn 325	Leu	Asp	Thr	Ile	Ile 330		Ser	Ala	Leu	Asp 335	Trp	
Glu Glu His			Arg	Phe	Gln									
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<pre>&lt;221&gt; CDS &lt;222&gt; (97). &lt;400&gt; 321 ggagttttt atcattgaga  aaa att tac Lys Ile Tyr  ttt ggg ggc Phe Gly Gly</pre>	ctggc tagtt aac Asn 10 aat Asn	ccc Pro cct Pro	aat Asn aca Thr	tct Ser agc Ser tgg	aca Thr atg Met 30	gaa Glu 15 ttt Phe	agt Ser gat Asp	atg Met 1 gtg Val ttg Leu	gaa Glu aat Asn aat Asn	gaa Glu aag Lys 35	ser aga Arg 20 atc Ile acc	cgc Arg 5 aag Lys aag Lys	aag Lys att Ile tat Tyr	114
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											atc Ile 100			402
											caa Gln			450
											gaa Glu			498
_			_			_	_		_		gat Asp	-		546
											gaa Glu			594
											gct Ala 180			642
		-				_			_		ttt Phe			690
											att Ile			738
				-	_	_		_			aac Asn	_		786
	_			_	_	 _			•	_	caa Gln	_		834
											gaa Glu 260			882
		-		_		 _					act Thr		-	930
-					_			_	_	_	ttg Leu	~	•	978
											aaa Lys			1026

260 265 270

Leu Gly Leu Thr Ser Ser Leu Ile Glu Gln Tyr Ile Gln Phe Leu Ala
275 280 285

Asp Ser Arg Leu Ser Lys Val Gly Ile Ala Lys Val Tyr Gly Val Gln

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His Pro Ile Lys Trp Val Glu Ser Phe Ser Ser Phe Asn Glu Gln Arg
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Ser Phe Asp Asp Phe
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tttttgttgg attctaac atg aaa gat ttt tat ttc atg gag atg aac act
                    Met Lys Asp Phe Tyr Phe Met Glu Met Asn Thr
cgt ttg caa gtg gaa cac acc att agc gaa atg gtg agc ggg tta aac
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Arg Leu Gln Val Glu His Thr Ile Ser Glu Met Val Ser Gly Leu Asn
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                                  20
                                                      25
ctc att gag tgg atg att aaa atc gct caa ggc gaa aaa ttg ccc aag
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Leu Ile Glu Trp Met Ile Lys Ile Ala Gln Gly Glu Lys Leu Pro Lys
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                             35
caa gaa agc ttt tct ctc aaa ggg cat gcg ata gaa tgc cga atc acg
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Gln Glu Ser Phe Ser Leu Lys Gly His Ala Ile Glu Cys Arg Ile Thr
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gca gaa gat cct aaa aaa ttc tac cca agc ccg ggc aaa att acc gaa
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Ala Glu Asp Pro Lys Lys Phe Tyr Pro Ser Pro Gly Lys Ile Thr Glu
                     65
tgg atc gct cct ggt ggg gtg aat gtg cgc ctt gat tcg cac gcg cat
                                                                   411
Trp Ile Ala Pro Gly Gly Val Asn Val Arg Leu Asp Ser His Ala His
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                                      85
gcc aat tat gtc gtg cct acg cac tat gat tcg atg att ggc aag ctc
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Ala Asn Tyr Val Val Pro Thr His Tyr Asp Ser Met Ile Gly Lys Leu
             95
                                100
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Ile Val Trp Gly Glu Asn Arg Glu Arg Ala Ile Ala Lys Met Lys Arg
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Ala Leu Lys Glu Phe Lys Val Glu Gly Ile Lys Thr Thr Ile Pro Phe
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                    145
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                            40
                                                 45
Leu Lys Gly His Ala Ile Glu Cys Arg Ile Thr Ala Glu Asp Pro Lys
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Lys Phe Tyr Pro Ser Pro Gly Lys Ile Thr Glu Trp Ile Ala Pro Gly
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                                         75
Gly Val Asn Val Arg Leu Asp Ser His Ala His Ala Asn Tyr Val Val
Pro Thr His Tyr Asp Ser Met Ile Gly Lys Leu Ile Val Trp Gly Glu
                                105
Asn Arg Glu Arg Ala Ile Ala Lys Met Lys Arg Ala Leu Lys Glu Phe
                            120
Lys Val Glu Gly Ile Lys Thr Thr Ile Pro Phe His Leu Glu Met Leu
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		-		-			_		_	tgc Cys	_		344
 		_								agc Ser		_	392
					_	_			_	agc Ser	_		440
										cat His			488
										gtg Val 160			536
										gtt Val			584
										caa Gln			632
										gtg Val			680
										tta Leu			728

	215	220		225
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	-		tct cat aaa ttc Ser His Lys Phe 255	
		-	ttg aat agt tat Leu Asn Ser Tyr 270	=
		_	ggg cgc tac acc Gly Arg Tyr Thr 285	
			cgt ttt atc acg Arg Phe Ile Thr	
		_	aaa aac gag caa Lys Asn Glu Gln 320	~
	Thr Arg Ala		acc ttg ctg tgc Thr Leu Leu Cys 335	
			agc gtg aat atg Ser Val Asn Met 350	
			tat tgc cat ttt Tyr Cys His Phe 365	_
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			gaa tta gag agc Glu Leu Glu Ser 400	
	Ile Ala Ile	-	gat cac act agc Asp His Thr Ser 415	•
			aac gct caa aaa Asn Ala Gln Lys 430	
atc tta atc ggc	act caa atg	ata agc aaa	ggg cat gat tac	gct aaa 1400

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agt t Ser T																1496
ggg a Gly A		_	_													1544
gaa a Glu T 5																1592
caa t Gln T 515			_		-		_	_			-					1640
ctg t Leu C																1688
ttg a Leu S										-	-		-	_		1736
gta a Val T							_			_			-			1784
tat c Tyr A 5																1832
aaa a Lys S 595																1880
gtg a Val A									taaa	aaac	etc a	ıtgtt	ttat	a		1927
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Ser Thr Glu Thr Asp Leu Leu Glu Asn Phe Leu Glu Asp Tyr Glu Asp
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Gln Gln Leu Ser Leu Lys Ala Ser Gln Thr Leu Ser Ser Cys Leu Glu
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_		_		_			ttg Leu			_		_	_	~	591
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	_			-	-		aat Asn	_			_		_		687
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				gcc Ala								1119
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				cac His								1551
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			gtg Val													1791
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	_	-	tgg Trp 590	-					_	_	_	_			_	1887
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			cag Gln	-	-							-				1983
		_	aaa Lys							_	-	_	-	-		2031
-	-		tac Tyr	-			_	_					_		_	2079
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			att Ile													2175
	_	_	gct Ala						_						_	2223
			gaa Gl _. u													2271

atc o																2319
gtg ( Val (																2367
aaa ( Lys (		taat	tata	aag (	gataa	atato	gt ga	ataga	agcti	t ga	catta	aacg	cta	gcga	taa	2423
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Gln V		Asp	Val	Gly	Lys 70		Val	Leu	Gln	Ile 75		Cys	Gly	Ala	Lys 80	
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His (	Gly	Met 115	Ile	Cys	Ser	Ser	Ile 120	Glu	Leu	Gly	Phe	Pro 125	Lys	Ile	Asn	
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Gly V	/al	Ile	Met 260		Ala	Tyr	Ser	Leu 265		Thr	Thr	Pro	Met 270		Leu	
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Lys Gln Thr Glu His Ser Leu Lys Asp Arg Thr Leu Thr Phe Gln Leu
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Gln Gly Ile Leu Lys Asn Leu Gly Phe Lys Val Ser Val Lys Glu Pro
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Gly Ile Asp Asn Leu Val Ser Lys Pro Leu His Cys Val Ser Ser Lys
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Asn Ser Ile Ala Leu Ser Val Arg Cys Val Ile His Ser Leu Glu Lys

730

735

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ata ggg tat gag aat gtg agc ggg ttt gct ttt ggc atg ggg att gaa Ile Gly Tyr Glu Asn Val Ser Gly Phe Ala Phe Gly Met Gly Ile Glu 145 150 155	962
aga tta gcc atg ctg act tgc cag atc aat gat ttg cgc agt ttc ttt Arg Leu Ala Met Leu Thr Cys Gln Ile Asn Asp Leu Arg Ser Phe Phe 160 165 170	1010
gaa act gat ttg aga gtg ttg gag agc ttt taatgaaact gagcattaat Glu Thr Asp Leu Arg Val Leu Glu Ser Phe 175 180	1060
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His Tyr Phe Phe Gly Gly Val Lys Leu Arg Trp Arg Ser Ser Phe Phe 85 90 95  Pro Phe Thr Glu Pro Ser Ala Glu Val Asp Ile Ser Cys Val Phe Cys 100 110 110 110 110 110 110 110 110 11	
His Tyr Phe Phe Gly Gly Val Lys Leu Arg Trp Arg Ser Ser Phe Phe 85 90 95  Pro Phe Thr Glu Pro Ser Ala Glu Val Asp Ile Ser Cys Val Phe Cys 100 105 110 110 110 110 110 110 110 110	
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														aag Lys		200
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														gtt Val		296
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														agt Ser		392
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Leu Lys Ile Lys Glu Asn Asn Pro Pro Leu Ile Asp Glu Met Leu
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Tyr Phe Leu Ala Phe Phe Leu Val Gly Ala Tyr Gln Glu Val Leu Gly
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Glu Val Leu Lys Glu Ile Leu Ser Ser Tyr Gln Lys Arg Ala Leu Lys
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	_					gat Asp		-			_			_	_	536
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1				5					10					15		
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Ile Lys Gly Gln Ile Leu Leu Cys Ala Gln Gly His Asp Ala Thr Asn
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Asn Ala Arg Glu Arg Ile Lys Val Gly Gly Leu Pro Thr Phe Lys Phe
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Gly Ala Lys Lys Asn Ala Lys Glu Glu Gln Tyr Lys Gln Asp Glu Arg
Leu Asn Glu Arg Leu Arg Glu Phe Ala Glu Thr Leu Ile Glu Ser Val
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Arg Lys Leu Gln Lys Leu Gly Asp Tyr Glu Asn Ile Glu Lys Ile
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                            Met Gln Glu Arg Val Phe Lys Arg Lys
gtt tta gat gcg aat atc tta aaa gaa atg cat gcg aac aat gtc tgt
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Val Leu Asp Ala Asn Ile Leu Lys Glu Met His Ala Asn Asn Val Cys
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tat tcc aag cat tca aaa gat agg ttt att cct ttc aaa ttt gat aaa
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Tyr Ser Lys His Ser Lys Asp Arg Phe Ile Pro Phe Lys Phe Asp Lys
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Phe Gly Tyr Val Gly Cys Lys Leu Phe Lys Lys Ile Leu Asn Phe Pro
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                                 50
age aat aca act tte ttt ggt gge aca ggt tgt aag aaa ete atg gaa
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Ser Asn Thr Thr Phe Phe Gly Gly Thr Gly Cys Lys Leu Met Glu
ctt tta agt gaa atc gtt ata gat tct aga agt tct aaa att gcg tta
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25 Thr Glu Asn Gln Lys Lys Ala Ile Lys Ile Ala Leu Asn Thr Pro Asp

Leu	Leu 75	Ser	Glu	Ile	Val	Ile 80	Asp	Ser	Arg	Ser	Ser 85	Lys	Ile	Ala	Leu	
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													act Thr			448
								_		_		-	ttc Phe 135			496
	_			-				_				_	aat Asn			544
	-						_			-			ctc Leu		_	592
			-	_	-				-		_		ggt Gly	_		640
													aac Asn			688
													tat Tyr 215			736
		_			_								aag Lys	_		784
							_				_	_	tta Leu	_	_	832
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Gly Thr Gly Cys Lys Lys Leu Met Glu Leu Leu Ser Glu Ile Val Ile
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Asp Ser Arg Ser Ser Lys Ile Ala Leu Asn Arg His Tyr Ala Leu Thr
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Glu Arg Ile Gly Phe Leu Thr Ala Phe Lys Asn Lys Lys Gly Tyr Ile
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Gly Leu Asn Gly Glu Ser Asn Leu Pro Asn Ser Phe Phe Leu Gly Ile
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Cys Gly Tyr Leu Lys Lys Leu Phe Lys Lys Leu Lys Asp Arg Ala Phe
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Arg Leu Ala Asn Lys His Gly Val Phe Phe Leu Lys Ile Pro Lys His
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                                                    190
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Asn Asn Pro Cys Phe Ser Tyr Arg Leu Thr Tyr Asp Gln Leu Val Gly
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                                            220
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                                                                   104
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Ser	Phe 20	Leu	Arg	Ser	Ile	Ala 25	Phe	Phe	Asp	Asp	Phe 30	Ser	Ala	Ser	Phe	
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					aag Lys											296
	-				aaa Lys	_			_	_	_		_		-	344
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					cgc Arg 200											680
					atg Met											728
					ttt Phe											776

						tta Leu										824
						gaa Glu 265										872
						cag Gln										920
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	_		_			gct Ala	-							_	_	1016
_			_			ctt Leu		_		_	_					1064
			_			gct Ala 345	_		_		_	_				1112
						aac Asn										1160
						cgc Arg										1208
	_	-				aat Asn		_		_	-		_			1256
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						ttt Phe 425										1352
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						ggc Gly										1448

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											aaa Lys					1640
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											cta Leu	-		-	_	1784
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675	680	685	690
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Leu	Glu 900	Asn	Leu	Met	Ala	Gln 905	Thr	Ser	Leu	Glu	Gln 910	Gln	Lys	Asp	Phe	
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					ttg Leu											2888
					ttg Leu											2936
					aaa Lys											2984
					cta Leu				-							3032
	Leu				cgc Arg 1000	Leu					Asp					3080
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Asn	Ile	Tyr	Arg	Lys 85	His	Leu	Lys	Met	Pro 90	Leu	Glu	Lys	Ala	Arg 95	Glu	
Phe			T		ת 1 ת	T.011	Asp	Len		Lvc	Ara	Pro	Glv		Sor	
	Asp	Ala	100	Cys	AIA	пец		105	7114	my 5	,, <u>,</u>		110	9	Ser	

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His Phe Asp Lys His Tyr Lys Arg Ile Phe Lys Ala His Asp Trp Leu
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Glu Cys Leu Tyr Gln Ala Gln Phe Tyr Thr Ser Lys Asn Ile Glu Pro
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Phe Pro Ile Lys Asp Leu Ser Phe Glu Lys Met Arg Leu Lys Ala Lys
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Tyr Leu Leu Asp Asp Asn Ile Lys Asn Phe Glu Phe Ala Lys Ile Lys
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His Lys Ile Ser Leu Tyr Phe Lys Asp Ala Lys Phe Tyr Pro Lys Ser
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Val Ala Leu Gly Phe Ser Ser Leu Phe Glu Asn Lys Leu Lys Lys Asn
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Lys Gly Arg Glu Ser Phe Ala Leu Pro Leu Ile Leu Asn Glu Glu Lys
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Lys Arg Gly Lys Ser Ser Asp Leu Phe Glu Trp Ala Leu Glu Gln Leu
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Lys Leu Glu Cys Gly Gly Ile Ser Ile Ile Gly Glu Asp Arg Asn Asn
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Ser Arg Gln Cys Lys Gly Ser Val Asn Lys Asp Glu Leu Ser Leu Gly
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Ala Arg Val Cys Leu Glu Val Gly Pro Asp Lys Asn Asp His Gly Lys
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Tyr Arg Gly Lys Ile Tyr Gly Leu Glu Lys Asn Arg Glu Ile Val Leu
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Leu Asn Thr Ala Lys Asn Ser Tyr Gln Arg Lys Pro Leu Asp Glu Lys
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Ile Lys His Arg Ile Glu Ala Leu Lys Arg Ile Lys Tyr Pro Cys Leu
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Lys Ile Phe Ser His Tyr Met Leu Glu Glu Leu Glu Thr Leu Asn Pro
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Tyr Phe Asp Pro Gln Thr Asp Arg Asp Phe Lys Lys Gly Leu Leu Asp
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Phe Phe Ser Arg Leu Asn Asp Ser Ile Pro Ala Lys Leu Gln Gln Glu
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Phe Ile Asn Leu Pro Ser Thr Asp Phe Leu Ser Arg Cys Leu Gly Ser
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Leu Glu Lys Asp Phe Gln Lys Thr Ile Phe Lys Lys Leu Lys Val Thr
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Asp Phe Leu Lys Arg Ile Glu Glu Cys Leu Lys Asn Pro Glu Ser Phe
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Tyr Phe Ser Ser Ala Cys Glu Leu Leu Leu Ala Phe Leu Ser Tyr Arg
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Arg Leu Leu Asp Ser Ile Asp Lys Ala Ile Glu Lys Glu Thr Glu Ile
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Lys Ser Phe Val Lys Leu Glu Leu Lys Asn Gln Ser Phe Asn Asn Ile
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Ser Phe Val Ile Lys Met Lys Phe Phe Lys Glu Lys Glu Val
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Ser Lys Ile Lys Ser Leu Arg Lys Phe Glu Ser Asn Pro Leu Val Arg
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Phe Asp Pro Ser Ala Leu Ala Leu Glu Pro Lys Phe
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Asn	Lys	Lys	Ala 150	Ile	Lys	Ala	His	Lys 155	Thr	Thr	Cys	Lys	Ser 160	cca Pro	Lys	536
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Met Pro

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Arg Ile Val Arg Ile Lys Ser Met Gly Met Ala Ala Ile Ser Trp Lys
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                              105
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Glu Ser Ser Lys Asn Phe Ser Thr Ile Ala Val Glu Glu Ser Glu Ser
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Pro Met Pro Lys Thr Arg Glu Lys Met Gly Glu Arg Pro Lys Thr Lys
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Tyr Pro Asn Lys Lys Ala Ile Lys Ala His Lys Thr Thr Cys Lys Ser
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Pro Lys Thr Ser Thr Ser Cys Leu Met Asp Leu Ser Leu Ser Lys Leu
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              165
Asn Ser Met Pro Ile Ile Asn Ile Lys Lys Thr Ile Pro Asn Ser Pro
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caa gga aaa ttt ttt tta cgc att gaa gac acg gat ttg agc cgt aac
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Gln Gly Lys Phe Phe Leu Arg Ile Glu Asp Thr Asp Leu Ser Arg Asn
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				ggt Gly										537
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				aga Arg 200										681
				aag Lys	-	_								729
		-	_	ttg Leu		-	-				_		-	777
				gtg Val										825
				ttt Phe				_		 		-	-	873
Lys				atg Met					Glu		_		Lys	921
275				280					285				290	

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ttg tta gaa Leu Leu Glu							
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aag gga ggc Lys Gly Gly 435	Gly Ile G		-	-		Leu Gly	
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Ala Asn Gln		Phe Phe I		Ile Glu	Asp Thr		Ser

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Tyr Tyr Cys Tyr Met Ser Lys Glu Glu Leu Asp Ala Leu Arg Glu Glu
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Gln Lys Ala Arg Lys Glu Thr Pro Arg Tyr Asp Asn Arg Tyr Arg Asp
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Phe Lys Gly Thr Pro Pro Lys Gly Ile Glu Pro Val Val Arg Ile Lys
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Val Lys Val Asn Thr Asn Glu Leu Asp Asp Phe Ile Ile Ala Arg Ser
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Asp Gly Thr Pro Thr Tyr Asn Phe Val Val Thr Ile Asp Asp Ala Leu
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Met Gly Ile Thr Asp Val Ile Arg Gly Asp Asp His Leu Ser Asn Thr
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Pro Lys Gln Ile Val Leu Tyr Lys Ala Leu Asn Phe Lys Ile Pro Asn
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Phe Phe His Val Pro Met Ile Leu Asn Glu Glu Gly Gln Lys Leu Ser
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Lys Arg His Gly Ala Thr Asn Val Met Asp Tyr Gln Glu Met Gly Tyr
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Leu Lys Glu Ala Leu Val Asn Phe Leu Ala Arg Leu Gly Trp Ser Tyr
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Gln Asp Lys Glu Val Phe Ser Met Gln Glu Leu Leu Glu Leu Phe Asp
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Pro Lys Asp Leu Asn Ser Ser Pro Ser Cys Phe Ser Trp His Lys Leu
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Asn Trp Leu Asn Ala His Tyr Leu Lys Asn Gln Ser Val Gln Glu Leu
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Pro Thr Gln Leu Asp Arg Leu Leu Asp Ala Leu Lys Glu Arg Ser Gln
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Thr Leu Lys Glu Leu Ala Leu Lys Ile Asp Glu Val Leu Ile Ala Pro
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Val Glu Tyr Glu Glu Lys Val Phe Lys Lys Leu Asn Gln Ala Leu Val
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Met Pro Leu Glu Lys Phe Lys Leu Glu Leu Asn Lys Ala Asn Phe
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Asn Asp Glu Ser Ala Leu Glu Asn Ala Met Arg Gln Ile Ile Glu Glu
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Glu Lys Ile Lys Ala Gly Ser Phe Met Gln Pro Leu Arg Leu Ala Leu
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Lys Gly Ile Tyr Pro Met Arg Asn Phe Pro Ile His His Asn Gly Phe
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Lys His Glu Val Leu Ala His Met Leu Lys Arg His Lys Glu Pro Phe
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Ile Leu Ser Tyr Asn Asp Cys Glu Phe Val Arg Asn Ala Tyr Lys Asp
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Phe Lys Ile Leu Glu Pro Ser Trp Gln Tyr Thr Met. Gly Gln Gly Glu
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aaa caa tct cat gag tta ttg att atc aag gag taaaaatgca tattagcgaa 349
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Lys Asp Phe Lys Ile Leu Glu Pro Ser Trp Gln Tyr Thr Met Gly Gln
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Gly Glu Ile Arg Met Gly Lys Asn Arg Leu Glu Arg Gly Asp Asn Asn
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ataggggtga ttatagcatt tatggggcaa aaaaagtaga atctgtatca agttttatta 180
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Ala Asp Ala Thr Thr Leu Leu Gly Phe Phe Glu Glu Asn Gln Asn Asn
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Gln Phe Val Ile Pro Ile Tyr Gln Arg Leu Tyr Ser Trp Lys Lys Glu
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gcc tac cag cat Ala Tyr Gln His 275	_	•		_	920
ctc tat ata gag Leu Tyr Ile Glu	-	•	l Ser Ile Ile		968
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Glu Arg Tyr Met Arg Phe Tyr Asn Pro Asn Phe Thr Arg Tyr Asp Gly 210 215 220	
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										gaa Glu				672
				Ala						cct Pro				720
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	tct Ser														816
	acg Thr														864
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	aaa Lys														960
	aaa Lys														1008
	gcc Ala	-		_	_		_		_	_	_	_	-		1056
	agc Ser							-		-	_	-	_	_	1104
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	tgc Cys				-	_	_			_					1200
	aca Thr														1248
	gca Ala														1296
	ata Ile														1344
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act g																2304
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acc as Thr Ly 8!																2592
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Gln Ala Ser Glu Cys Leu Asp Asn Leu Asp Asp Pro Thr Asp Gln Glu Ala Ile Glu Gln Cys Leu Glu Gly Leu Ser Asp Ser Glu Arg Ala Leu Ile Leu Gly Ile Lys Arg Gln Ala Asp Glu Val Asp Leu Ile Tyr Ser Asp Leu Arg Asn Arg Lys Thr Phe Asp Asn Met Ala Ala Lys Gly Tyr Pro Leu Leu Pro Met Asp Phe Lys Asn Gly Gly Asp Ile Ala Thr Ile Asn Ala Thr Asn Val Asp Ala Asp Lys Ile Ala Ser Asp Asn Pro Ile Tyr Ala Ser Ile Glu Pro Asp Ile Ala Lys Gln Tyr Glu Thr Glu Lys 520. Thr Ile Lys Asp Lys Asn Leu Glu Ala Lys Leu Ala Lys Ala Leu Gly Gly Asn Lys Lys Asp Asp Asp Lys Glu Lys Ser Lys Lys Ser Thr Ala 55Û Glu Ala Lys Ala Glu Asn Asn Lys Ile Asp Lys Asp Val Ala Glu Thr Ala Lys Asn Ile Ser Glu Ile Ala Leu Lys Asn Lys Lys Glu Lys Ser Gly Glu Phe Val Asp Glu Asn Gly Asn Pro Ile Asp Asp Lys Lys Lys Ala Glu Lys Gln Asp Glu Thr Ser Pro Val Lys Gln Ala Phe Ile Gly Lys Ser Asp Pro Thr Phe Val Leu Ala Gln Tyr Thr Pro Ile Glu Ile Thr Leu Thr Ser Lys Val Asp Ala Thr Leu Thr Gly Ile Val Ser Gly Val Val Ala Lys Asp Val Trp Asn Met Asn Gly Thr Met Ile Leu Leu Asp Lys Gly Thr Lys Val Tyr Gly Asn Tyr Gln Ser Val Lys Gly Gly Thr Pro Ile Met Thr Arg Leu Met Ile Val Phe Thr Lys Ala Ile Thr Pro Asp Gly Val Ile Ile Pro Leu Ala Asn Ala Gln Ala Ala Gly Met Leu Gly Glu Ala Gly Val Asp Gly Tyr Val Asn Asn His Phe Met Lys Arg Ile Gly Phe Ala Val Ile Ala Ser Val Val Asn Ser Phe Leu Gln Thr Ala Pro Ile Ile Ala Leu Asp Lys Leu Ile Gly Leu Gly Lys Gly Arg Ser Glu Arg Thr Pro Glu Phe Asn Tyr Ala Leu Gly Gln Ala Ile Asn Gly Ser Met Gln Ser Ser Ala Gln Met Ser Asn Gln Ile Leu Gly Gln Leu Met Asn Ile Pro Pro Ser Phe Tyr Lys Asn Glu Gly Asp Ser Ile Lys Ile Leu Thr Met Asp Asp Ile Asp Phe Ser Gly Val Tyr Asp Val Lys Ile Thr Asn Lys Ser Val Val Asp Glu Ile Ile Lys Gln Ser 

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     Met Ala Val Tyr Leu Asp Phe Glu Asn His Ile Lys Glu Ile Gln
aat gaa att gaa tta gcc ctt att aga ggc gat gag gac gct aaa gaa
                                                                   157
Asn Glu Ile Glu Leu Ala Leu Ile Arg Gly Asp Glu Asp Ala Lys Glu
atc tta gaa aaa aga ttg gat aag gag gtt aaa agc att tat tcc aat
                                                                   205
Ile Leu Glu Lys Arg Leu Asp Lys Glu Val Lys Ser Ile Tyr Ser Asn
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                                                      45
ctc act gat ttt caa aaa ctc caa tta gca aga cac cct gac aga ccc
                                                                   253
Leu Thr Asp Phe Gln Lys Leu Gln Leu Ala Arg His Pro Asp Arg Pro
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tac gct atg gat tac att gat ctc atc tta aaa gat aaa tat gaa gtc
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Tyr Ala Met Asp Tyr Ile Asp Leu Ile Leu Lys Asp Lys Tyr Glu Val
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ttt ggg gat agg cat tat aac gat gat aaa gcg atc gtg tgc ttt gta
                                                                   349
Phe Gly Asp Arg His Tyr Asn Asp Asp Lys Ala Ile Val Cys Phe Val
ggg aaa att gat aat gtc cca gtt gtg gtg atc gga gaa gaa aag ggc
                                                                   397
Gly Lys Ile Asp Asn Val Pro Val Val Val Ile Gly Glu Glu Lys Gly
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aga ggg act aaa aac aaa ctc tta aga aat ttt ggc atg cct aac cct
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Arg Gly Thr Lys Asn Lys Leu Leu Arg Asn Phe Gly Met Pro Asn Pro
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                                                     125
tgt ggc tat cgt aag gct ttg aaa atg gca aag ttt gct gaa aag ttt
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Cys Gly Tyr Arg Lys Ala Leu Lys Met Ala Lys Phe Ala Glu Lys Phe
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                                                 140
aat ttg cct att tta atg ctt gtg gat aca gcc ggg gcg tat ccg ggg
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Thr Asp Phe Gln Lys Leu Gln Leu Ala Arg His Pro Asp Arg Pro Tyr 50 55 60

Ala Met Asp Tyr Ile Asp Leu Ile Leu Lys Asp Lys Tyr Glu Val Phe 65 70 75 80

Gly Asp Arg His Tyr Asn Asp Asp Lys Ala Ile Val Cys Phe Val Gly 85 90 95

Lys Ile Asp Asn Val Pro Val Val Val Ile Gly Glu Glu Lys Gly Arg
100 105 110

Gly Thr Lys Asn Lys Leu Leu Arg Asn Phe Gly Met Pro Asn Pro Cys 115 120 125

Gly Tyr Arg Lys Ala Leu Lys Met Ala Lys Phe Ala Glu Lys Phe Asn 130 135 140

Leu Pro Ile Leu Met Leu Val Asp Thr Ala Gly Ala Tyr Pro Gly Ile 145 150 155 160

Gly Ala Glu Glu Arg Gly Gln Ser Glu Ala Ile Ala Lys Asn Leu Gln
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       Met Gln Phe Thr Gly Lys Asn Val Leu Ile Thr Gly Ala Ser
aaa ggc att ggg gct gaa atc gcc aaa act ctc gct tct atg ggg ctg
                                                                   156
Lys Gly Ile Gly Ala Glu Ile Ala Lys Thr Leu Ala Ser Met Gly Leu
 15
                     20
aaa gtt tgg atc aat tac cgc agt aat gct gaa gtg gct gac gct ttg
                                                                   204
Lys Val Trp Ile Asn Tyr Arg Ser Asn Ala Glu Val Ala Asp Ala Leu
                 35
aaa aat gag ctt gaa gaa aaa ggc tat aag gca gct gtc att aaa ttt
                                                                   252
Lys Asn Glu Leu Glu Glu Lys Gly Tyr Lys Ala Ala Val Ile Lys Phe
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                                 55
                                                      60
                                                                   300
gat gcg gct tct gaa agc gat ttt att gaa gcg ata caa acc atc gtc
Asp Ala Ala Ser Glu Ser Asp Phe Ile Glu Ala Ile Gln Thr Ile Val
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caa agc gat ggg ggg ttg tct tac ttg gtg aat aac gcc ggt gtg gtg
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Gln Ser Asp Gly Gly Leu Ser Tyr Leu Val Asn Asn Ala Gly Val Val
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cgc gat aaa tta gcg atc aaa atg aaa aca gaa gac ttt cac cat gtc
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Arg Asp Lys Leu Ala Ile Lys Met Lys Thr Glu Asp Phe His His Val
ata gac aat aac ctc act tca gcc ttt ata ggt tgc cga gag gct tta
                                                                   444
Ile Asp Asn Asn Leu Thr Ser Ala Phe Ile Gly Cys Arg Glu Ala Leu
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aag gtg atg agc aag agt cgt ttt ggg agc gtg gtc aat gtc gct tct
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Lys Val Met Ser Lys Ser Arg Phe Gly Ser Val Val Asn Val Ala Ser
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                                135
atc att ggt gaa aga ggc aat atg ggg cag aca aac tac tca gcg agt
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Ile Ile Gly Glu Arg Gly Asn Met Gly Gln Thr Asn Tyr Ser Ala Ser
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Lys				Ile					Ser							
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				aac Asn					_	_						732
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мес 1	PLO	αтλ	тте	∟ys 5	val	AT G	GIU	στλ	Asp 10	AIG	rue	ASP	GIU	Ala 15	ıyr	

Arg	Arg	Pne	цуs 20	гур	GIII	1111	ASP	25	ASII	пец	vai	Val	30	GIU	Cys	
Arg	Ala	Arg 35	Arg	Phe	Phe	Glu	Ser 40	Lys	Thr	Glu	Lys	Arg 45	Lys	Lys	Gln	
Lys	Ile 50	Ser	Ala	Lys	Lys	Lys 55	Val	Leu	Lys	Arg	Leu 60	Tyr	Met	Leu	Arg	
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